1 A NanoLuc luciferase-based assay enabling the real-time analysis of protein secretion

- 2 and injection by bacterial type III secretion systems
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- 13 Running Head: NanoLuc-based T3SS secretion and injection assay
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- 16
- 17 Word count abstract: 160
- 18 Word count text: 6669

19 Abstract

The elucidation of the molecular mechanisms of secretion through bacterial protein secretion 20 systems is impeded by a lack of assays to quantitatively assess secretion kinetics. Also the 21 analysis of the biological role of these secretion systems as well as the identification of 22 inhibitors targeting these systems would greatly benefit from the availability of a simple, quick 23 and quantitative assay to monitor principle secretion and injection into host cells. Here we 24 present a versatile solution to this need, utilizing the small and very bright NanoLuc luciferase 25 to assess secretion and injection through the type III secretion system encoded by Salmonella 26 pathogenicity island 1. The NanoLuc-based secretion assay features a very high signal-to-noise 27 ratio and sensitivity down to the nanoliter scale. The assay enables monitoring of secretion 28 kinetics and is adaptable to a high throughput screening format in 384-well microplates. We 29 further developed NanoLuc and split-NanoLuc-based assays that enable the monitoring of type 30 31 III secretion-dependent injection of effector proteins into host cells.

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33 Importance

The ability to secrete proteins to the bacterial cell surface, to the extracellular environment, or 34 35 even into target cells is one of the foundations of interbacterial as well as pathogen-host interaction. While great progress has been made in elucidating assembly and structure of 36 37 secretion systems, our understanding of their secretion mechanism often lags behind, not last because of the challenge to quantitatively assess secretion function. Here, we developed a 38 39 luciferase-based assay to enable the simple, quick, quantitative, and high throughputcompatible assessment of secretion and injection through virulence-associated type III secretion 40 systems. The assay allows detection of minute amounts of secreted substrate proteins either in 41 the supernatant of the bacterial culture or within eukaryotic host cells. It thus provides an 42 enabling technology to elucidate the mechanisms of secretion and injection of type III secretion 43 systems and is likely adaptable to assay secretion through other bacterial secretion systems. 44

45 Introduction

The ability to secrete proteins to the bacterial cell surface, to the extracellular environment, or 46 even into target cells is one of the foundations of interbacterial as well as pathogen-host 47 interaction. Protein export is particularly challenging for Gram-negative bacteria as two 48 membranes of the bacterial cell envelope have to be passed. So far, nine different protein 49 secretion systems, named type I – IX secretion systems (T1SS – T9SS), have been discovered 50 in Gram-negative bacteria (1, 2). Three of these systems, T3SS, T4SS, and T6SS, serve the 51 direct application of effector proteins into target cells of either prokaryotic or eukaryotic origin 52 53 (3).

54 Due to its form and function, the type III secretion machine, as used by many enteric pathogens like Salmonella, Shigella, Yersinia, or enteropathogenic Escherichia coli, is called injectisome 55 (4). It is composed of a base that anchors the machine to the inner and outer membranes of the 56 bacterial cell envelope (5), of cytoplasmic components that serve in targeting and receiving of 57 58 substrates (6, 7), of an inner membrane-localized export apparatus performing substrate 59 unfolding and export (8), and of a needle filament through which secreted substrates reach the 60 host cell (9) (Fig. 1A). Injection itself is mediated by a needle tip complex and by hydrophobic translocators forming pores in the host cell's target membrane (10). Type III secretion is 61 62 energized by ATP hydrolysis of the system's ATPase and by the proton motive force (PMF) 63 across the bacterial inner membrane (11). Secretion of substrates follows a strict hierarchy with early substrates building up the needle filament, intermediate substrates forming the needle tip 64 and translocon pore, and late substrates that serve as effectors inside the target cell. 65

While great progress has been made in elucidating assembly and structure of the type III 66 secretion injectisome (12-14), our understanding of its secretion mechanism lags behind, not 67 68 last because of the challenge to quantitatively assess secretion function. Traditionally, T3SS function is assessed by SDS PAGE, Western blotting, and immunodetection of secreted 69 70 substrates, either acid precipitated from the bacterial culture supernatant, or analyzed in lysates of eukaryotic target cells (15). This approach is time-consuming, at best semi-quantitative, and 71 72 lacks sensitivity. To facilitate a simplified analysis of principle secretion, injection, and intracellular localization, several enzyme-linked and fluorescent reporters have been developed 73 74 (16).

Ampicillin resistance conferred by β-lactamase-fusions secreted into the periplasm was used to
monitor the function of flagellar T3SS, which are closely related to T3SS of injectisomes (17).
Secretion into the periplasm through partially assembled injectisomes was assessed by using

PhoA-fusions, instead (18). While these assays proved very valuable to address some specific 78 questions, monitoring of secretion into the periplasm is only sensible for early substrates as 79 switching to the secretion of later substrates does not occur without an assembled needle. High 80 throughput (HTP) assays for screening of T3SS inhibitors exploited the turn-over of the 81 fluorogenic substrate PED6 by a secreted phospholipase fusion (19), the turn-over of the 82 chromogenic cephalosporine nitrocefin by a secreted β -lactamase fusion (20), and the 83 enzymatic uncaging of the fluorogenic substrate Glu-CyFur by a secreted carboxypeptidase 84 85 fusion (21, 22).

Likewise, several reporter assays have been developed to assess the injection of T3SS effectors 86 into eukaryotic host cells. Pioneering work by the Cornelis lab exploited the specific increase 87 in intracellular cAMP levels upon injection of effectors fused to a calmodulin-activated 88 adenylate cyclase (Cya) (23). Later, this assay was also adapted to assay injection of effectors 89 by T4SS (24). While the Cya assay showed a very good signal to noise ratio (S/N) of several 90 logs, it was not suitable to monitor injection kinetics or to be adapted for HTP screening because 91 92 of a tedious cAMP analysis protocol. Widely used to assay injection of effector proteins in T3SS 93 and T4SS is an assay that utilizes the enzymatic cleavage of the FRET-reporter cephalosporin CCF2 by injected β-lactamase-fusions (25). The CCF2 assay facilitated the analysis of injection 94 kinetics and of intracellular accumulation levels of effectors (26). It was also successfully used 95 for HTP high content screening of T3SS inhibitors (27). Real-time observation of injection was 96 achieved by direct fluorescent labeling of tetracysteine motif-tagged effectors (28). However, 97 since this approach requires multidimensional time-lapse microscopy, it is not feasible for 98 routine analysis of effector injection or HTP. Split-GFP technology (29) and self-labelling 99 enzyme tags (30) were successfully used to monitor intracellular localization of effector 100 proteins but both techniques are not optimal for the analysis of translocation kinetics: split-GFP 101 because of a low sensitivity and the slow kinetics of GFP complementation, and the self-102 labelling enzyme tags because labelling can only be done with effectors that have already been 103 104 translated before host cell contact.

We aimed to develop a T3SS assay based on effector-luciferase fusions to enable a simple, quantitative, and HTP-compatible assessment of principle secretion and injection. The advantage of luciferase-reporters is a very high S/N and sensitivity. In addition, luciferase-based assays benefit from the lack of product (light) accumulation, simplifying the analysis of secretion and injection rates. We developed a secretion assay utilizing NanoLuc (NLuc) luciferase, an engineered 19 kDa glow-type luciferase from the deep-sea shrimp *Oplophorus* gracilirostris that converts furimazine, emitting blue light (31). The NLuc-based secretion assay

allowed quantification of minute amounts of secreted effectors either in the supernatant of the bacterial culture or within eukaryotic host cells. The assay's ultra-high sensitivity, its wide dynamic range and quick response dynamics qualify it as an enabling technology to elucidate the mechanisms of secretion and injection of T3SS and is likely adaptable to assay secretion through other bacterial secretion systems.

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118 Results

119 Assessment of effector-luciferase fusion proteins as reporters for type III secretion

In order to identify a luciferase compatible with type III secretion through the T3SS encoded 120 121 by Salmonella pathogenicity island 1 (SPI-1, T3SS-1), we evaluated six different commercially available luciferases as effector-fused secretion reporters: Cypridinia luciferase (CLuc), 122 Gaussia princeps luciferase (GLuc), Gaussia dura luciferase (GDLuc), NLuc, Renilla luciferase 123 (RLuc), and Red Firefly luciferase (RFLuc) (31-35). We generated translational fusions of the 124 effectors SipA and SopE, respectively, coupled at their C-termini to a luciferase and a myc 125 epitope-tag. The effector-luciferase fusions were expressed from a rhamnose-inducible pro-126 moter on a low-copy number plasmid in wild type S. Typhimurium and in a secretion deficient 127 mutant ($\Delta sctV$). The expression and type III-dependent secretion of the effector luciferase fu-128 sions was assessed by SDS PAGE, Western blotting and immunodetection of the myc epitope 129 tag in whole bacterial cells and culture supernatants, respectively, after 5 h of growth. All ef-130 131 fector-luciferase fusions could be detected at the expected molecular mass in whole cells and in culture supernatants, indicating their productive expression and secretion (Fig. 1A). CLuc 132 and RFLuc showed additional bands likely corresponding to the cleaved luciferase-myc. In 133 134 general, SipA-luciferase fusions were secreted more efficiently than SopE fusions. SipA and SopE fusion with CLuc as well as SopE fusions with NLuc and RLuc could only be detected in 135 very low levels in the culture supernatants (Fig.1B). 136

137 The activity of the secreted luciferases in filtered culture supernatants of the *S*. Typhimurium 138 wild type and of the $\Delta sctV$ mutant, respectively, was assessed by luminometry using the speci-139 fied conditions for each luciferase. The S/N (wild type vs. $\Delta sctV$) was highest with effector-

140 NLuc fusions (SipA-NLuc S/N = 45, SopE-NLuc S/N = 22), and, with the exception of GDLuc,

141 always higher for SipA-luciferase fusions (Fig. 1C).

142 Since the SipA-NLuc fusion showed the best S/N, we introduced SipA-NLuc-myc into the

143 chromosome of a S. Typhimurium wild type strain and of a $\Delta sctV$ mutant for further analysis.

144 First, we compared the expression and secretion of plasmid and chromosome-encoded SipA-

145 NLuc, respectively, and as a reference also of the secreted translocator SctE, by SDS PAGE,

146 Western blotting and immunodetection. SipA-NLuc was expressed well from the chromosome

147 even though, not unexpectedly, at lower levels compared to its expression from the plasmid

- 148 (Fig. 1D). The extent of T3SS-dependent secretion of plasmid and chromosome-encoded SipA-
- 149 NLuc was indistinguishable (Fig. 1D).

We next evaluated the S/N of the secreted SipA-NLuc fusion when expressed from plasmid or chromosome by measuring the NLuc activity in filtered culture supernatants of the wild type and the $\Delta sctV$ mutant. While plasmid-based expression resulted in a S/N = 45, chromosomebased expression even achieved a S/N = 200. The stronger plasmid-based expression may lead

to a greater liberation of SipA-NLuc upon occasional cell lysis, compromising the S/N.

Both, injectisomes and flagella possess T3SS for the export of proteins and it has been shown 155 that substrates of one system may be secreted by the other one to a limited degree (36, 37). In 156 order to assess the contribution of the flagellar T3SS to the S/N of SipA-NLuc secretion, we 157 158 blocked expression of flagella by deleting the gene of the flagellar master regulator FlhD. In the absence of flagella, the S/N of SipA-NLuc secretion increased to 140 when SipA-NLuc was 159 160 expressed from the plasmid and to 1000 when it was expressed from the chromosome (Fig. 1E). FlhD contributes to the induction of the SPI-1-encoded T3SS by indirectly regulating the ex-161 pression of the the major SPI-1 regulator HilA (38), which results in a strongly decreased ex-162 163 pression of T3SS-1 and its effectors in the absence of FlhD. To determine whether the improved S/N of SipA-NLuc secretion in the *flhD* mutant resulted from an overall lower expression of 164 the reporter or from preventing secretion through flagella, we also tested SipA-NLuc secretion 165 in a strain expressing chromosome-encoded HilA from an arabinose-inducible promoter (39), 166 thus uncoupling its expression from control by FlhD. In this strain, T3SS-1-dependent SipA-167 NLuc secretion was identical in the wild type and in the *flhD* mutant (Fig. 1F). However, in the 168 absence of a functional T3SS-1 (AsctDFLJ), 150-fold lower levels of SipA-NLuc were detecta-169 ble in the culture supernatant of the strain lacking flagella. These results indicate that about 1% 170 of the SipA-NLuc secretion signal in the wild type strain stems from secretion through the 171 flagellar T3SS (Fig. 1E) and that the increased S/N in the absence of FlhD results from prevent-172 173 ing secretion through flagella. Despite the increased S/N in the absence of flagella, we used *flhD* wild type bacteria for most of the work presented herein because of the higher overall 174 175 signal and because motility appeared to promote growth in a microplate format.

In order to test the versatility of NLuc as secretion reporter, we also constructed fusions withthe early T3SS substrate SctP (needle length regulator) and with the intermediate substrate SctA

(tip protein). While NLuc compromised secretion and function of SctP when fused to its C-178 terminus (Fig. S1AB), SctA-NLuc fusions were readily secreted, even when NLuc was placed 179 at different positions within the polypeptide chain of SctA (Fig. S1CD). To overcome the limi-180 tation of NLuc in supporting secretion of SctP, we utilized a split-NLuc approach. Split-Nluc is 181 composed of a large fragment (LgBiT, 18 kDa) comprising most of the protein's beta barrel and 182 of a small fragment with a high affinity to the LgBiT (HiBiT, 1.3 kDa), comprising only one 183 beta strand (40). SctP-HiBiT fusions were successfully secreted into the culture supernatant and 184 strong luminescence was detected when complementing SctP-HiBiT with LgBiT (Fig. S1AB), 185 186 showing that split-NLuc can serve as a secretion reporter when NLuc fails.

In summary, we could show that luciferases are versatile reporters for T3SS and that effectorNLuc fusions report on secretion with a very high S/N, even in the absence of plasmid-based
overexpression.

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191 Assessment of the sensitivity of the NLuc-based secretion assay

One handicap of the traditional, Western blot-based secretion assay is its low sensitivity that impedes analyzing low culture volumes as required for the analysis of secretion kinetics or for the development of HTP screens.

In order to compare the sensitivity of the Western blot- and the SipA-NLuc-based secretion assays, we made a serial dilution of the filtered supernatant of wild type and $\Delta sctV S$. Typhimurium cultures grown for 5 h. In the Western blot-based assay, we could detect the intermediate substrate SctE down to a supernatant volume of 113 µl and the early substrate SctP as well as the late substrate SipA-NLuc down to 225 µl (Fig. 2A). In contrast, using the SipA-NLuc assay, we were able to obtain a stable S/N = 200 down to 195 nl supernatant volume. The S/N even remained above 50 when assaying an equivalent of only 24 nl (Fig. 2B).

Next, we assessed the performance of the SipA-NLuc assay in monitoring the onset kinetics of 202 type III secretion, which requires very high sensitivity due to the small amounts of secreted 203 material that is initially present. To this end, we grew S. Typhimurium harboring arabinose-204 controlled HilA to an $A_{600} = 0.9$, after which expression of the pathogenicity island was induced 205 by the addition of 0.02% (w/v) arabinose. Bacterial cells and culture supernatants were col-206 lected every 10 min and kept on ice until reading at the end of the experiment. Induction of SPI-207 1 was monitored by Western blot and immunodetection of the base component SctJ in whole 208 209 cells. It was first observed 30 min after the addition of arabinose (Fig. 2C). Also luminescence of SipA-NLuc was detected in the culture supernatant for the first time 30 min after induction 210

of *S*PI-1 and then luminescence increased steadily to the end of the measurement after 120 min (Fig. 2C). This increase in luminescence correlates directly with SipA-NLuc secretion and is not influenced by NLuc maturation or turn-over as the activity of NLuc remains stable in the culture supernatant over extended periods of time (Fig. S2).

Both experiments, serial dilution and secretion kinetics, prove the superior sensitivity of the NLuc-based over the traditional secretion assay. While the detection of secreted substrate proteins using the traditional assay requires larger volumes and accumulation of substrates in the culture supernatant for an extended period of time, the NLuc assay allows detection of secretion in very small volumes, in brief intervals, and with very short handling times (10 min after collection of supernatant). Our results also show that induction and assembly of the megadalton injectisome is a very quick process that gets bacteria rapidly armed for attack.

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223 Application example: Harnessing the NLuc secretion assay for high throughput screening

The high sensitivity and the short handling time of the SipA-NLuc-based secretion assay provided an excellent basis to develop a HTP assay for drug screening in a 384-well microplate format.

Centrifugation or filtering is not feasible for separation of bacterial cells and culture supernatant 227 228 in a microplate format. In order to overcome this problem, we made use of the high-protein binding capacity of the microplates and tested whether secreted substrates would specifically 229 230 bind to the plate wall after being secreted (Fig. 3A). To this end, 50 µl of S. Typhimurium wild type and $\Delta sctV$ mutant were grown in white high protein binding 384-well plates. Bacteria were 231 232 washed out of the wells after 5 h of growth using a microplate washer. Then, PBS, NLuc buffer, and NLuc substrate were supplied to each well and the luciferase activity was measured. Using 233 234 this setup, a S/N = 37 could be achieved (Z' = 0.8), which is excellent for HTP screening (Fig. 235 3B).

To assess the robustness of this assay and the variation across the plate, we filled an entire 384-236 well plate with 50 µl of a S. Typhimurium, SipA-NLuc culture and allowed it to grow for 5 h at 237 37°C. Luminescence of secreted, wall-bound SipA-NLuc was assessed after washing out bac-238 teria as described above. The assay proved very robust with a coefficient of variation of 7% 239 over the entire plate and with little edge effects (Fig. 3C, Table S1). We then performed a proof-240 of-concept inhibitor screen by assessing the effect of a range of 37 different bioactive com-241 pounds on the activity of the T3SS in the 384-well format (Table S2, Fig. 3D). Each well of the 242 plate was printed with 0.5 µl of a compound in 100% DMSO, to which 50 µl of a S. 243

Typhimurium, SipA-NLuc culture was added. Again, the culture was allowed to grow for 5 h, 244 after which secretion of SipA-NLuc was assessed by luminometry. The assay showed a highly 245 dynamic response from 10 % to 120 % secretion activity compared to the DMSO-treated wild 246 type control (Fig. 3D). Detection of SipA-NLuc was most strongly reduced by the flavonoids 247 quercetin (30 µg/ml, 90% reduction) and scutellarin (10 µg/ml, 75% reduction), which con-248 firms the previously reported observation that flavonoids target T3SS (22). Also treatment with 249 the 3-hydroxy-3-methylglutaryl (HMG) coenzyme A reductase-blocker simvastatin reduced de-250 tection of SipA-NLuc by 44%. Replication of the screen proved a high reproducibility of the 251 assay with a R^2 of 0.95 (Fig. 3E). 252

Over all, the SipA-NLuc assay proved to be highly adaptable to a high throughput screening format in 384-well plates, featuring a high S/N, a low error across the plate, a great reproducibility and requiring only short hands-on time.

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257 Application example: Assessment of the PMF-dependence of type III secretion by the NLuc
258 secretion assay

It has been known for long that secretion through T3SS depends on two sources of energy, on 259 the hydrolysis of ATP by the system's ATPase (FliI in flagella, SctN in injectisomes) and on the 260 PMF (41-43), which itself is composed of the ΔpH , i.e., the proton concentration gradient across 261 the membrane, and the $\Delta \Psi$, the electric potential difference between the periplasm and cyto-262 263 plasm. The contribution of these two PMF components to T3SS function can be dissected with 264 specific inhibitors. Carbonyl cyanide 3-chlorophenylhydrazone (CCCP) is a PMF uncoupler (ionophore) and discharges both the ΔpH and the $\Delta \Psi$ by transporting protons through the mem-265 266 brane (44). At acidic pH, potassium benzoate is a weak acid and can enter the membrane and discharge the ΔpH (45). Valinomycin can shuttle potassium ions across the membrane which 267 268 collapses the electric potential difference $\Delta \Psi$ (46). Evaluating the contribution of each PMF component to T3SS function requires the careful analysis of secretion kinetics, for which the 269 270 classical, semi-quantitative Western blot-based secretion assay is not well suited, but for which the NLuc-based secretion assay proved very powerful. To further show this, CCCP, potassium 271 272 benzoate, and valinomycin, respectively, were added to the bacterial culture at different concentrations, 60 min after induction of SPI-1 (for experimental details, please refer to the meth-273 ods section), while samples of culture supernatants were taken every 10 min for subsequent 274 analysis of the luminescence of secreted SipA-NLuc. While SipA-NLuc secretion progressed 275 over time in the control sample (Fig. 4), addition of the inhibitors lead to sudden changes in 276

secretion kinetics. CCCP blocked secretion instantly, even at concentrations of 5 μ M, showing 277 the critical relevance of the PMF for type III secretion (Fig. 4A). Discharching the ΔpH by 278 potassium benzoate resulted in a concentration-dependent instant reduction of secretion (Fig. 279 4B). At 20 mM potassium benzoate, secretion was completely abolished while it proceeded at 280 60% of the untreated control in the presence of 5 mM and at 10% in the presence of 10 mM 281 potassium benzoate. Collapsing the electric potential by valinomycin lead to a strongly reduced 282 luciferase signal after 10 min, after which secretion proceeded in a concentration-dependent 283 manner (Fig. 4C): in the presence of 20 µM valinomycin, no significant change in secretion 284 285 rate was observed, while 40 μ M and 60 μ M valinomycin, respectively, lead to 70% and 40% secretion of the untreated control. 286

These results show that both components of the PMF, ΔpH and $\Delta \Psi$, contribute to energizing secretion in the *S*PI-1-encoded T3SS of *S*. Typhimurium. As the PMF-compromising compounds took effect so quickly after treatment, it is highly unlikely that the PMF-dependence of type III secretion is the consequence of a secondary effect of PMF reduction – an issue that could only be resolved with the sensitive and highly time-resolved NLuc secretion assay. These results open the door for further experiments dissecting the role of the different T3SS components in utilizing the PMF.

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295 Development of NLuc-based host cell injection assays

Assessment of secretion of T3SS substrates into the culture supernatant is very useful for investigating the basic secretion mechanism of T3SS, however the intended biological function of T3SS injectisomes is the injection of effector proteins into host cells. Since the SipA-NLucbased secretion assay proved to be very sensitive and simple, we attempted to adapt the assay to monitoring the injection of SipA-NLuc into host cells.

301 In a first and simple approach, we infected HeLa cells in 96-well plates at an MOI = 50 with SipA-NLuc-expressing S. Typhimurium, using wild type bacteria and secretion-deficient $\Delta sctV$ 302 mutants. After infection for 60 min, attached bacteria were gently washed off with PBS using a 303 microplate washer and subsequently, the HeLa cell-associated luminescence was measured us-304 ing live cell buffer (Fig. 5A). The non-secreting $\Delta sctV$ mutants (Fig. 5A) showed a HeLa cell-305 associated luminescence of 8% of the wild type, corresponding to a S/N = 12 (Fig. 5C). To 306 307 determine whether the HeLa cell-associated signal was truly resulting from injected SipA-NLuc, we assessed injection in a set of mutants that are capable of secreting SipA but incapable 308

309 of injecting it into host cells: a needle tip-deficient $\Delta sctA$, a translocon-deficient $\Delta sctEBA$, and

a gatekeeper-deficient $\Delta sctW$ mutant. While secretion of SipA-NLuc into the culture superna-310 tant was increased between 2 and 5-fold in $\triangle sctA$, $\triangle sctEBA$, and $\triangle sctW$ mutants (Fig. 5B), 311 which are reportedly unlocked for secretion of late substrates like SipA (47, 48), the HeLa cell-312 associated luminescence was strongly reduced to 9-24% of the wild type when infecting with 313 these mutants (Fig. 5C). From these results we can conclude that the luminescence signal ob-314 tained from infection with wild type S. Typhimurium resulted to more than 90% from injected 315 SipA-NLuc and that only little signal may stem from bacteria remaining attached to HeLa cells 316 or to the plate even after washing. Over all, this NLuc-based injection assay proved very useful 317 318 for the quick and simple assessment of translocation of effectors into host cells by an end-point measurement, however the kinetics of injection cannot be assessed by this assay. 319

To gain a higher specificity for the signal of injected SipA and enable analysis of injection 320 kinetics, we employed the split version of the NLuc luciferase. To this end, SipA was fused to 321 322 HiBiT while LgBiT was expressed stably by the HeLa cell line. Complementation of LgBiT 323 with HiBiT to a functional luciferase should only occur inside the HeLa cells after translocation of SipA-HiBiT (Fig. 5D). We first tested the secretion of SipA-HiBiT into the culture superna-324 tant by providing LgBiT to the assay buffer. Similar to what was observed for SipA-NLuc, 325 326 secretion of SipA-HiBiT into the culture supernatant was increased between 2 and 6-fold in $\Delta sctA$, $\Delta sctEBA$, and $\Delta sctW$ mutants, respectively (Fig. 5E). However, in contrast to the SipA-327 NLuc-based injection assay, none of the T3SS mutant strains yielded any detectable lumines-328 cence in the split NLuc assay (Fig. 5F), making this assay highly suitable for monitoring the 329 specific injection of T3SS effectors into host cells. This setup even allowed us to follow the 330 kinetics of SipA-HiBiT injection over time directly in a microplate reader (Fig. 5G). 331

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333 Discussion

334 The elucidation of the molecular mechanisms of secretion through T3SS and other bacterial protein secretion systems is impeded by a lack of assays to quantitatively assess secretion 335 kinetics. Also the analysis of the biological role of these secretion systems as well as the 336 identification of inhibitors targeting these systems would greatly benefit from the availability 337 of a simple, quick and quantitative assay to monitor principle secretion and injection into host 338 cells. Here we present a versatile solution to this need, utilizing the small and very bright NLuc 339 luciferase to assess secretion and injection through the T3SS encoded by SPI-1 of S. 340 Typhimurium. Secretion of a SipA-NLuc fusion showed a very high S/N and sensitivity down 341 to the nanoliter scale, making it exquisitely suited for the assessment of secretion kinetics. In 342

addition, the NLuc-based secretion assay proved highly adaptable to a HTP screening format
in 384-well microplates. We further developed NLuc and split-NLuc-based assays that enable
the monitoring of T3SS-dependent injection of effector proteins into host cells.

A perfect assay to monitor protein secretion would feature: i) A lack of signal from the un-346 secreted reporter, resulting in a high S/N. ii) A small reporter that does not interfere with 347 secretion through the secretion system of interest. In case of T3SS, this also includes a not too 348 fast and tight folding inside bacteria as only unfolded protein can be secreted and as the 349 unfolding capacity of the system is not very high. iii) A fast and efficient folding of the reporter 350 outside of the bacterium, guaranteeing fast response dynamics. iv) An intrinsic signal of the 351 reporter, not necessitating an enzyme substrate. v) A high sensitivity. vi) A lack of accumulation 352 of product of the reporter's reaction. And vii) Be quick, simple, and needing only short hands-353 354 on time.

While fluorescent proteins would be desirable secretion reporters as they benefit from an 355 356 intrinsic signal (and thus do not come with the problem of accumulation of product of the reporter's reaction), they often suffer from a very slow maturation time and/or insufficient 357 brightness. In addition, fluorescent proteins tend to form very stable β-barrels that block 358 secretion through T3SS (49), excluding them as secretion reporters, at least for T3SS. While 359 the use of split GFP can overcome the limitation associated with tight folding, slow 360 complementation and maturation of GFP compromise its use. The NLuc-based secretion assay 361 362 as presented herein matches most of the needs listed above. While NLuc lacks an intrinsic signal and requires the addition of a substrate, the analysis of secretion by this assay is not complicated 363 by the overlay of the kinetics of the reporting enzyme and the kinetics of secretion, as it is in 364 other enzyme-linked secretion assays. Instead, the measured signal of the NLuc assay is directly 365 proportional to the amount of accumulated secreted protein. This advantage, together with the 366 superior sensitivity, yield a very high dynamic range of the NLuc secretion assay. 367

We demonstrated that the NLuc-secretion assay is highly suited to study the kinetics of secretion due to its superior sensitivity. Our simple assay setup only allowed deduction of secretion kinetics from the accumulation of NLuc in the culture supernatant but culturing bacteria in a microfluidics system could enable the direct and on-line reading of secretion into the medium flow through and by this facilitate an even better resolved analysis of the mechanism of secretion.

Our experiments show that secretion of NLuc is supported by fusion to a range of intermediateand late T3SS substrates, even within a polypeptide chain, but fails to be secreted when fused

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to the early substrate SctP. It is conceivable that the mode of early substrate secretion does not 376 provide a sufficient unfolding capacity to support secretion of NLuc while this seems not a 377 problem when NLuc is fused to intermediate and late substrates. Interestingly, a Yersinia SctP-378 PhoA fusion was secreted (18), pointing either to a higher unfolding capacity of the Yersinia 379 T3SS or to a weaker fold of PhoA. We could overcome the limited use of NLuc as secretion 380 reporter for early substrates by using the split-NLuc technology instead. The 11 amino acid-381 long HiBiT was accommodated well by SctP and it is conceivable that this very small piece 382 allows secretion in most circumstances. 383

In its current form, the NLuc secretion assay requires the separation of bacteria and supernatant to achieve a good S/N because of the membrane-permeating properties of the NLuc substrate. A membrane impermeant NLuc substrate could overcome this limitation, would make NLucbased secretion assays even more simple and versatile and increase their robustness due to less steps of handling.

In addition to the points important for a secretion assay, a perfect injection assay would also: i) Feature a high specificity for injected effectors as opposed to secreted but not injected ones. ii) Allow the analysis of injection kinetics. And iii) Allow localization of the injected protein, at best at single molecule resolution.

While fluorescence-based assays proved highly suitable to study the localization dynamics of 393 injected effectors inside host cells, they are very limited in their use to study injection kinetics 394 and are always instrumentation-demanding. The CCF2-based injection assay features simple 395 396 handling, instead, and proves very useful for the analysis of injection, but suffers from high costs of CCF2 and a low dynamic range. In addition, the product accumulation resulting from 397 the enzymatic activity of the injected β -lactamase complicate the analysis of injection kinetics. 398 The herein-presented NLuc-based injection assays offer very quick and simple analysis, even 399 400 of injection kinetics, and feature a high dynamic range and sensitivity. While a high-resolution analysis of the localization of the effector-NLuc-fusions inside host cells is not supported by 401 402 these assays, microscopic setups exist that utilize luminescence for long-duration monitoring of single cells (50), which may become useful for studying the role of individual effectors in 403 bacterial persistence. 404

As performed herein, cytoplasmic expression of LgBiT will only generate luminescence if the HiBiT of the injected effector also localizes to the cytoplasm. However, the split-NLuc injection assay may also be utilized to analyze the localization and topogenesis of effector proteins inside host cells by targeting LgBiT to specific organelles instead (Fig. 6). Furthermore,

complementation of LgBiT by the low-affinity SmBiT instead of the high-affinity HiBiT may
provide a useful tool to investigate effector-host protein interactions *in vivo* by bimolecular
complementation (51).

In summary, our data show that NLuc-fusions of secreted substrate proteins can be used as a robust, versatile, cheap, simple and quick reporter for T3SS secretion and injection that will enable future in-depth elucidation of T3SS function (Fig. 6). The NLuc reporter is likely to be adaptable to other bacterial secretion systems as well.

416

417 Materials and methods

418 *Materials*

Chemicals were from Sigma-Aldrich unless otherwise specified. SERVAGel[™] TG PRiME[™]
8–16% precast gels were from Serva. Primers, listed in Table S3, were synthetized by Eurofins
and Integrated DNA Technologies. Monoclonal anti-c-myc antibody was from Roche (11-667149-001). Secondary antibodies goat anti-mouse IgG DyLight 800 conjugate were from
Thermo-Fisher (SA5-35571).

424

425 Bacterial strains, plasmids and growth conditions

Bacterial strains and plasmids used in this study are listed in Table S3. All Salmonella strains 426 were derived from Salmonella enterica serovar Typhimurium strain SL1344 (Hoiseth and 427 Stocker, 1981) and created by allelic exchange as previously described (52). S. Typhimurium 428 strains were cultured with low aeration at 37°C in Lennox broth (LB) supplemented with 0.3 M 429 NaCl to induce expression of SPI-1. As required, bacterial cultures were supplemented with 430 tetracycline (12.5 µg/ml), streptomycin (50 µg/ml), or kanamycin (25 µg/ml). Plasmids were 431 432 generated by Gibson cloning (53) using KOD (Novagen) or Q5 polymerase (NEB). Expression of pT10-based plasmids was induced by the addition of 100 µM of rhamnose to the culture 433 434 medium.

435

436 Western-blot-based secretion assay

Western-blot-based analysis of type III-dependent secretion of proteins into the culture medium
was carried out as described previously (39). S. Typhimurium was cultured at 37°C for 5 h. For
separation of whole cells and cell culture supernatant, the bacterial suspensions were

440 centrifuged at $10,000 \times g$ for 2 min at 4°C. Whole cells were directly resuspended in SDS PAGE

loading buffer. The supernatant was filtered through a $0.22 \ \mu m$ pore size filter, sodium deoxy-

442 cholic acid was added to a final concentration of 0.1% (w/v), and proteins were precipitated by

443 addition of 10% trichloroacetic acid (v/v; final concentration) for 30 min at 4°C. After pelleting

444 by centrifugation at $20,000 \times g$ for 20 min at 4°C, precipitated proteins were washed with ace-

- tone and subsequently resuspended in SDS PAGE loading buffer.
- 446

447 *Luciferase assays*

To measure NLuc, RFLuc, Gluc, GDluc, Rluc and Cluc activity of secreted translational fu-448 sions, bacteria were grown under SPI-1-inducing conditions for 5 h. Culture supernatants were 449 separated from whole bacterial cells by centrifugation for 2 min at 10,000 x g. The following 450 buffers were prepared with their substrates according to the manufacturers' protocols: For Nluc, 451 25 µl of Nano-glo assay buffer containing furimazine (Nluc working solution, Promega) was 452 added to 25 µl of the culture supernatant. For RFLuc, 30 µl of constituted One-glo assay buffer 453 containing luciferin (Promega) was added to 30 µl of the culture supernatant. For Gluc and 454 GDLuc, 50 µl of the assay buffer containing coelenterazine (Thermo Fisher) was added to 20 455 µl of culture supernatant. For RLuc, 25 µl of the constituted assay buffer (Promega), in which 456 the substrate was 1:100 diluted, was added to 25 µl of the culture supernatant. For CLuc, a 457 working solution was prepared containing assay buffer and 1:100 of the substrate vargulin 458 (Thermo Fisher). 30 µl of the working solution was added to 10 µl of the supernatant. The 459 luciferase activities were measured in white 384-well plates (MaxiSorp, Nunc), with acquisition 460 settings as recommended by the manufacturers. 461

462

463 *NLuc assay for wall-bound protein*

In order to measure wall-bound protein, overnight cultures of *S*. Typhimurium were back-diluted to an $A_{600} = 0.1$ and 50 µl of the bacterial suspension was transferred to a 384-well microplate (MaxiSorp, Nunc) and grown at 37° for 5 h. The plate was washed with water using a microplate washer (Tecan Hydrospeed) and the Nluc working solution was diluted in PBS (30 µl PBS + 10 µl NLuc working solution) and added to each well to measure luminescence using the Tecan Spark reader with following settings: attenuation: auto, settle time: 0 ms, integration time: 100 ms.

- 471 For the inhibitor screen, 0.5 μl of each compound (Table S2) was added to 50 μl bacterial culture
- 472 prior to incubation at 37°C for 5 h, and the plate was processed as described above.
- 473

474 SDS PAGE, Western blotting and immunodetection

For protein detection, samples were separated by SDS PAGE using SERVAGelTM TG PrimeTM 8-16% precast gels and transferred to a PVDF membrane (Bio-Rad) by standard protocols. Membranes were probed with primary antibodies α -SctP (39), α -SctE (39), α -c-Myc and α -SctJ (39). Secondary antibodies were goat anti-mouse IgG DyLight 800 conjugate. Detection was performed using the Odyssey imaging system (Li-Cor).

480

481 MBP-NLuc and MBP-HiBiT expression and purification

NLuc and HiBiT, respectively, were cloned into a pMal-c5X vector to yield a translational fu-482 483 sion with maltose-binding protein (MBP). E. coli BL21 was transformed with the plasmids. Bacterial cultures were grown overnight at 37°C in LB broth and back-diluted in Terrific Broth 484 (TB) the next day to an $A_{600} = 0.1$. They were grown to an $A_{600} = 0.6-0.8$ at 37°C. Subsequently, 485 expression of MBP-NLuc/ HiBiT was induced by addition of IPTG to a final concentration of 486 487 0.5 mM, after which bacteria were further grown at 37°C for 4 h. Bacterial cells were harvested by centrifugation (6,000 x g, 15 min, 4° C) and resuspended in column binding buffer (CB) 488 containing 200 mM NaCl, 20 mM Tris-HCl (pH 7.4), 1 mM EDTA, Protease inhibitor (Sigma-489 Aldrich, P8849, 1:100), DNAse 10 µg/ml, 1 mM MgSO₄ and lysozyme (10 µg/ml) and lysed 490 with a French press. The obtained solution containing cell lysate and cell debris was two times 491 centrifuged at 15,000 x g for 20 min at 4°C. MBP-NLuc/HiBit in the clear lysate was bound to 492 an amylose resin (NEB), washed with CB and eluted by 10 mM maltose in the same buffer. 493 Buffer was exchanged to PBS by using the Amicon Ultra system (Merck). 494

495

496 *Stability test of NLuc*

497 40 μ l Purified MBP-NLuc was added (2 μ g, final concentration) to 1 ml LB/ 0.3 M NaCl and 498 to 1 ml culture supernatant of wild type *S*. Typhimurium. Samples were kept either at 37°C, at 499 room temperature, or on ice for up to 4 h. Aliquots were removed over time and transferred to 500 a 384-well plate, 25 μ l of the NLuc working solution was added and luminescence was directly 501 measured in a microplate reader (Tecan Spark).

502

503 Kinetic measurement

SipA-NLuc was introduced into the chromosome of S. Typhimurium, Para-hilA by allelic ex-504 change. The resulting strain was grown overnight at 37°C in LB/0.3 M NaCl, and was back-505 diluted the following day to an $A_{600} = 0.1$. Bacterial cultures grew to an $A_{600} = 0.9$ in an Erlen-506 meyer flask in a 37°C water bath, stirred with a magnet stirrer. Expression of SPI-1 was induced 507 by addition of arabinose to a final concentration of 0.02% (v/v) and samples were taken at 508 different time points thereafter for assessment of the luminescence of secreted SipA-NLuc or 509 for immunodetection of SctJ. For testing the role of PMF inhibitors, bacterial cells were washed 510 511 twice after reaching an $A_{600} = 0.9$ in LB/0.3 M NaCl containing either 120 mM Tris-HCl, pH 7.0 for CCCP (Sigma) or 120 mM Tris-HCl, pH 7.0 and 150 mM KCl for valinomycin (Sigma). 512 For potassium benzoate, cells at the same growth stage ($A_{600} = 0.9$) were harvested and then 513 washed twice with LB/0.3 M NaCl containing 80 mM MES buffer, pH 6.8. The cultures in the 514 515 different media (without inhibitor, with 0.02% (v/v) arabinose) were kept in the water bath at 37°C and 200 µl of samples were taken at different time points and kept on ice. The inhibitors 516 517 were added to the bacterial culture 60 min after *hilA*-induction. Cultures were kept in the water bath and samples were taken every 10 min. Samples were centrifuged to separate whole cells 518 519 and supernatant. 25 µl of the supernatant was transferred to a white 384-well plate and lumi-520 nescence was measured upon addition of the Nluc working solution in a luminometer.

521

522 *Generation of stable HeLa cell line expressing LgBiT*

LgBiT was cloned into the MCS of pLVX-EF1α-IRES-Puro (Takara) resulting in pLVX-EF1α-523 LgBiT-IRES-Puro by Gibson assembly. 24 h prior to transfection, three 10 cm cell culture plates 524 containing each 4 x 10⁶ HEK 293T cells in 8 ml DMEM supplemented with 10% FCS (v/v) 525 and sodium pyruvate were incubated at 37°C, 5% CO2 overnight. The next day, 7 µg DNA of 526 pLVX-EF1a-LgBiT-IRES-Puro in 600 µl sterile water was added to Lenti-X Packaging Single 527 Shot (Takara). The containing pellet was completely resuspended and the solution incubated 528 for 10 min at room temperature to allow formation of nanoparticle complexes. Finally the 529 DNA/nanoparticle solution was added dropwise to the HEK 293T cells. After 4 h of incubation 530 at 37°C, 6 ml growth medium was added and cell supernatant was harvested after 48 h and 531 sterile filtered. In total 42 ml supernatant were reduced to a total volume of 4.2 ml used Lentix-532 Concentrator (Takara) exactly according to the protocol of the manufacturer. The viral suspen-533 sion was aliquoted and stored at -80°C. The virus titer was determined using the QuickTiter 534

Lentivirus Titer Kit (Cell Biolabs) according to the manufacturers protocol. The viral superna-535 tant was then diluted to a final MOI of 2-10 in 10% FCS-VLE RPMI, supplemented with 536 polybrene (4 μ g/ml final concentration) and added to HeLa cells (5 x 10⁵ cell in 500 μ l medium 537 in six well plates). After overnight culture, medium was exchanged and cells were cultured for 538 another day. The cells were then split, transferred to cell culture plates, and 2 µg/ml puromycin 539 was supplemented. After outgrowth of stably transduced cells, single cell clones were generated 540 by single cell dilution. Various cell clones were tested and verified for LgBiT expression by 541 lysing the cells and performing a luciferase assay by the addition of purified MBP-HiBiT in the 542 543 Hibit Lytic Buffer from the Hibit Lytic Detection Kit. Buffer and substrate was added in 1:50 ratio as described in the manufacturer's protocol, MBP-Hibit (2 mg/ml) was added in 1:100 544 ratio to the buffer-substrate mixture. 545

546

547 Injection assay and injection kinetics

1 x 10⁴ HeLa cells and HeLa LgBiT cells were seeded out in white 96 well plates with glass 548 bottom 24 h before infection in 100 µl DMEM + 10% FCS (GIBCO). S. Typhimurium was 549 washed and resuspended in HBSS to infect the cells at a MOI = 50 for 60 min. After infection, 550 cells were gently washed with a microplate washer (Tecan Hydrospeed, 5 cycles dispensing and 551 aspirating (speed: 70 µl/sec)) using 1 x PBS (GIBCO). A final wash volume of 100 µl was 552 used together with 25 µl of Nanoglo live cell assay buffer (Promega) containing substrate for 553 luminescence measurement in a Tecan Spark reader with the following settings: attenuation: 554 auto, settle time: 0 ms, integration time: 1,000 ms. For monitoring the injection kinetics, HeLa 555 LgBit cells were seeded out and S. Typhimurium bacteria in HBSS were added to the cells as 556 described above. Directly upon addition of the bacteria, 25 µl of the reconstituted Nanoglo live 557 558 cell buffer was added to the infection culture and luminescence reading was carried out for 90 min with a 2 min reading interval in the Tecan Spark with the same settings as for the injection 559 560 assay.

561

562 Acknowledgements

We thank Thomas Hesterkamp and Mark Brönstrup for continued input in high throughput assay development. We acknowledge receipt of the LgBiT/HiBiT split luciferase system by Promega before commercial release. This work was funded in part by the German Center for Infection Research (DZIF), grant TTU06.801 WP1.

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- 700

701 Figure Legends

- **Fig 1** Assessing different luciferases as reporters for type III secretion.
- (A) Cartoon of the T3SS injectisome. Names or proteins mentioned herein are shown according
 to the unified nomenclature. The figure is adapted from reference (4).
- (B) Proteins of whole cell lysates and culture supernatants of *S*. Typhimurium expressing the
 indicated SipA-Luc and SopE-Luc fusions were analyzed by SDS PAGE, Western blot and
 Immunodetection with an anti-myc antibody.

708 (C) Signal to noise ratios (wt/ $\Delta sctV$) of luciferase activities of secreted SipA-Luc and SopE-

- Luc fusions were graphed. Bar graphs represent the mean S/N of three independentmeasurements.
- (D) Immunodetection of SipA-NLuc^{myc} and SctE on Western blot of SDS PAGE-separated
 culture supernatants and whole cell lysates, either expressing SipA-NLuc^{myc} from a
 plasmid or from the chromosome.
- 714 (E) Signal to noise ratios (wt/ $\Delta sctV$) of luciferase activities of secreted SipA-NLuc either 715 expressed from a plasmid or from the chromosome, each with or without flagella (*flhD*) 716 were graphed. Bar graphs represent the mean S/N of three independent measurements.
- 717(F) SipA-NLucmyc secretion in S. Typhimurium $P_{ara}hilA$ and in S. Typhimurium $\Delta sctDFIJ$,718 $P_{ara}hilA$ with and without flagella (*flhD*), respectively. Bar graphs represent mean (±719standard deviation) of three technical replicates. Asterisks indicate statistical significance720of SipA-NLucmyc secretion assessed by Student's *t*-test, *: $p \le 0.05$,

721 Abbreviations: Nluc: Nanoluc, RFLuc: Red Firefly luciferase, GDLuc: Gaussia Dura

- 722 Luciferase, GLuc: Gaussia princeps Luciferase, RLuc: Green Renilla Luciferase, CLuc:
- 723 Cypridinia Luciferase, S/N: signal to noise, ns: non-significant
- Fig 2 Assessment of the sensitivity of the NLuc secretion reporter
- (A) Immunodetection of the T3SS substrates SctP, SctE and SipA-NLuc^{myc} on a Western blot
 of SDS PAGE-separated, serially diluted culture supernatants.
- 727 (B) Luminescence of secreted SipA-NLuc^{myc} in serially diluted culture supernatants of the 728 S. Typhimurium wild type and a $\Delta sctV$ mutant. Triangles show the calculated signal to 729 noise ratios for each dilution. Data represent the mean (± standard deviation) of three 730 technical replicates.
- (C) Normalized luminescence of secreted SipA-NLuc^{myc} at different time points after induction
 of *hilA* with 0.02% arabinose. Experiments were normalized by setting the maximum
 luminescence of each experiment to 1. The data points represent mean (± standard
 deviation) of five independent measurements. At each time point, samples of whole cell
 lysates were taken for immunodetection of SctJ.
- **Fig 3** Development of a SipA-NLuc-based HTP secretion
- (A) Cartoon of the assay setup. *S*. Typhimurium expressing SipA-NLuc was grown in a 384 well microplate format. Secreted SipA-NLuc bound to the wall of the high protein binding microplate. Bacteria were washed out and luminescence was measured.
- (B) Luminescence and signal to noise ratio of secreted SipA-NLuc. The experimental setup
 was as shown in (A). Bars represent the mean (± standard deviation) of three technical
 replicates.
- 743 (C) Signal variation of SipA-NLuc secretion assayed over an entire 384-well microplate as
 744 shown in (A).
- (D) SipA-NLuc secretion in response to treatment with 37 different bioactive compounds,
 assayed as shown in (A). The DMSO-treated control was set to 100%. The layout of the
 plate is shown in Table S1.
- (E) Comparison of the results of two independent compound screens as in (D). The R² value
 was calculated from a linear regression.
- 750
- 751

23

752	Fig 4 Assessment of the PMF-dependence of type III secretion by the NLuc secretion assay
753	(A)Normalized secretion of SipA-NLuc in S. Typhimurium ParahilA after induction of SPI-
754	1 by addition of 0.02% arabinose. CCCP was added to a final concentration of 0, 5, 10
755	and 15 μ M, respectively, 60 min after induction of SPI-1.
756	(B) As in (A) but addition of K^+ benzoate to final concentration of 0, 5, 10 and 20 mM,
757	respectively.
758	(C) As in (A) but addition of Valinomycin to a final concentration of 0, 20, 40 and 60 μ M,
759	respectively.
760	All data represent means (\pm standard deviation) of three independent measurements.
761	
762	Fig 5 Development of NLuc-based host cell injection assays
763	(A) Cartoon showing setup of NLuc injection assay. S. Typhimurium expressing SipA-NLuc
764	was allowed to infect HeLa cells for 60 min. SipA-NLuc was injected into HeLa cells
765	by use of the T3SS injectisome. Bacteria were washed away using a microplate washer
766	and subsequently NLuc luminescence was measured.
767	(B) Luminescence of SipA-NLuc secreted by the S. Typhimurium wild type and indicated
768	mutants in the absence of host cells. The luminescence of the wild type was set to 100%.
769	(C) Luminescence of SipA-NLuc injected into HeLa cells by the S. Typhimurium wild type
770	and indicated mutants. The experimental setup was as shown in (A). The luminescence
771	of the wild type was set to 100%.
772	(D)Cartoon showing setup of split-NLuc (HiBiT) injection assay. S. Typhimurium
773	expressing SipA-HiBiT was allowed to infect HeLa cells (expressing LgBiT) for
774	60 min. SipA-HiBiT was injected into HeLa cells by use of the T3SS injectisome.
775	Luminescence of the complemented split-NLuc was measured.
776	(E) Luminescence of LgBiT-complemented SipA-HiBiT secreted by the S. Typhimurium
777	wild type and indicated mutants in the absence of host cells. The luminescence of the
778	wild type was set to 100%.
779	(F) Luminescence of SipA-HiBiT injected into LgBiT-expressing HeLa cells by the S.
780	Typhimurium wild type and indicated mutants. The experimental setup was as shown in
781	(D). The luminescence of the wild type was set to 100%.

782(G) Luminescence of SipA-HiBiT injected into LgBiT-expressing HeLa cells by the S.783Typhimurium wild type and the $\Delta sctV$ mutant. At timepoint zero, HeLa cells were784infected with S. Typhimurium after which cells were incubated inside a microplate785reader in the presence of NLuc substrate. Luminescence was followed in 2 min intervals.786Values of the $\Delta sctV$ mutant were set to zero for each time point. The results show the787mean of technical triplicates.

Bar graphs represent mean (\pm standard deviation) of three independent measurements. Asterisks indicate statistical significance between wt and mutant strains assessed by a Students *t*-test, ***: $p \le 0.001$ **: $p \le 0.01$

791

Fig 6 Cartoon summarizing the utilization of the NLuc-based T3SS secretion and injectionassays

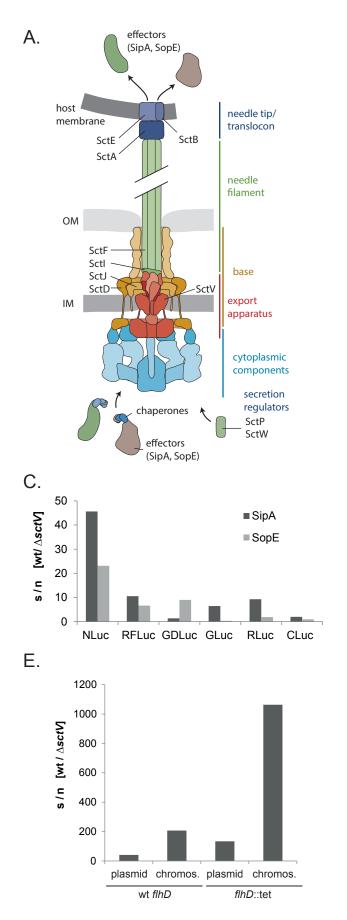
794

795 Supplemental Material

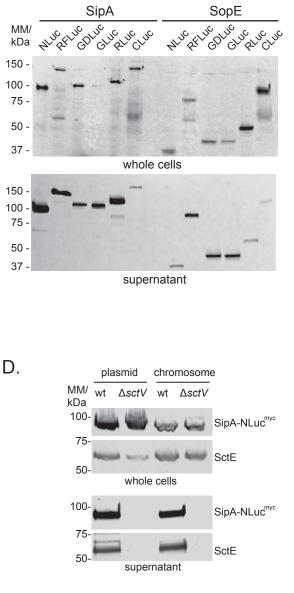
796 Fig S1 Expression and secretion of SctP-NLuc, SctP-HiBiT, and SctA-NLuc fusions

Fig S2 Stability of NLuc in LB and in culture supernatant

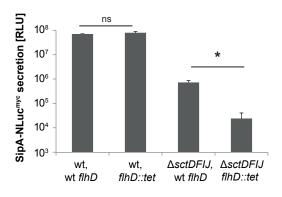
- **Table S1** Statistics of the reproducibility assessment of the 384-well microplate format NLuc secretionassay
- 800 Table S2 Layout of compound screening test plate incl. SipA-NLuc secretion of one measurement
- 801 Table S3 Strains, plasmids, primers







F.



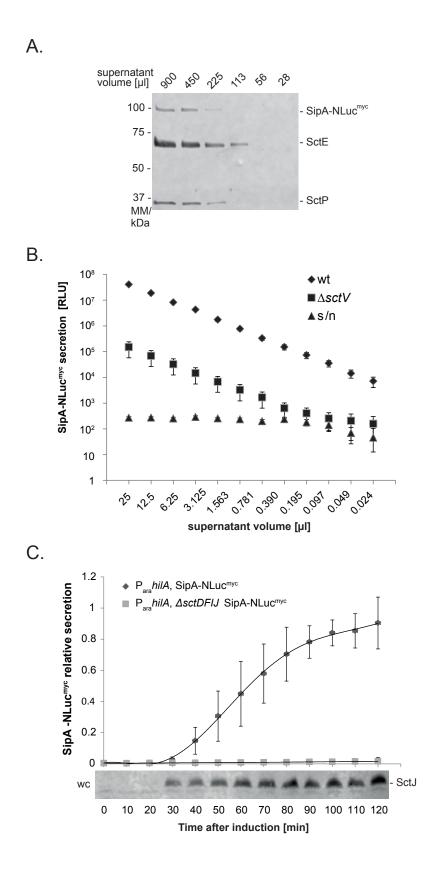
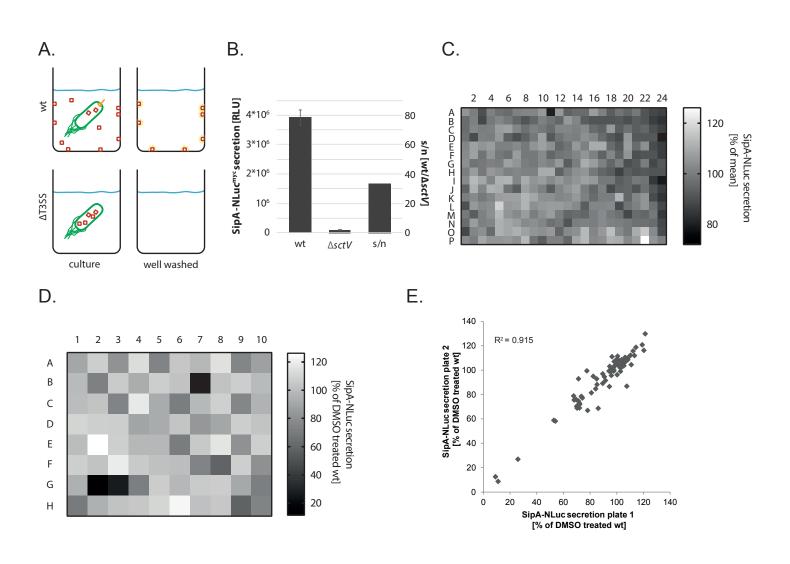


Figure 2



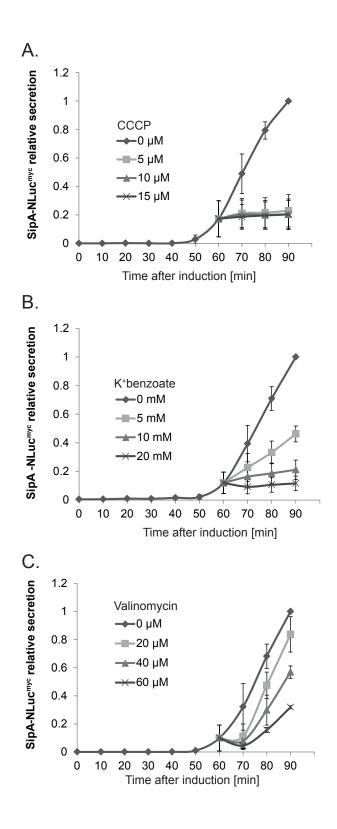
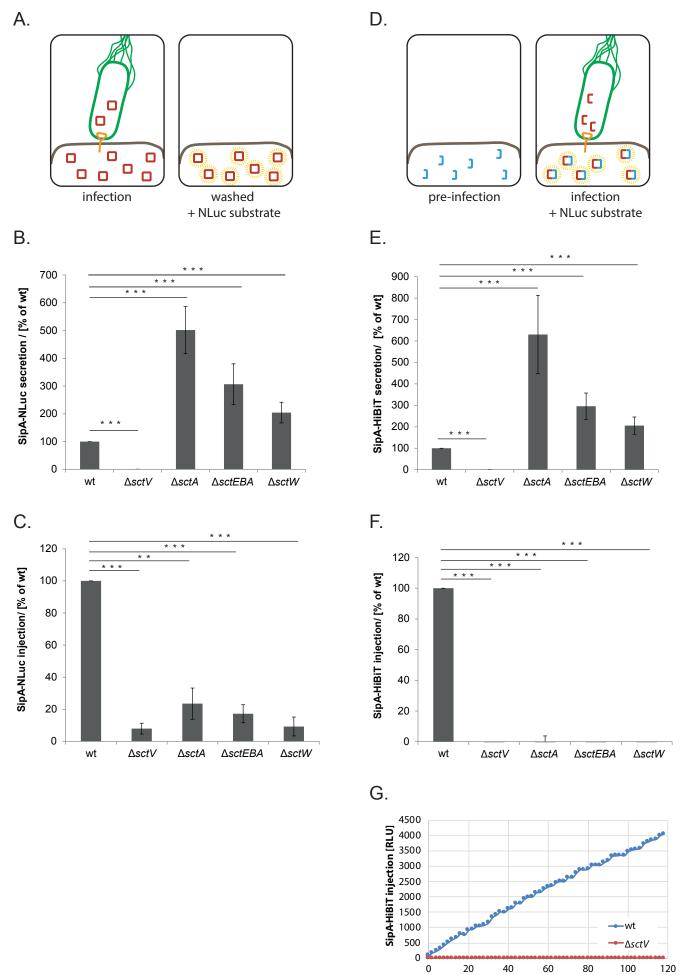


Figure 4



time [min]

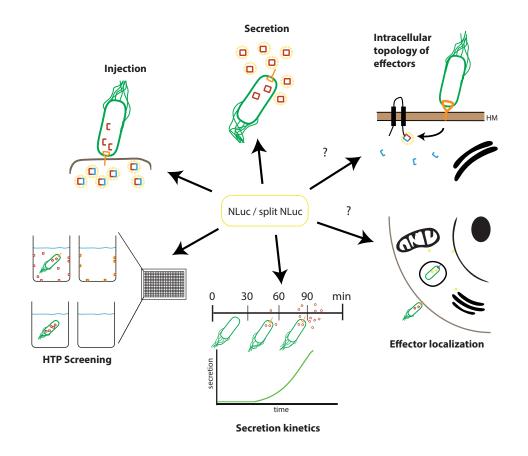
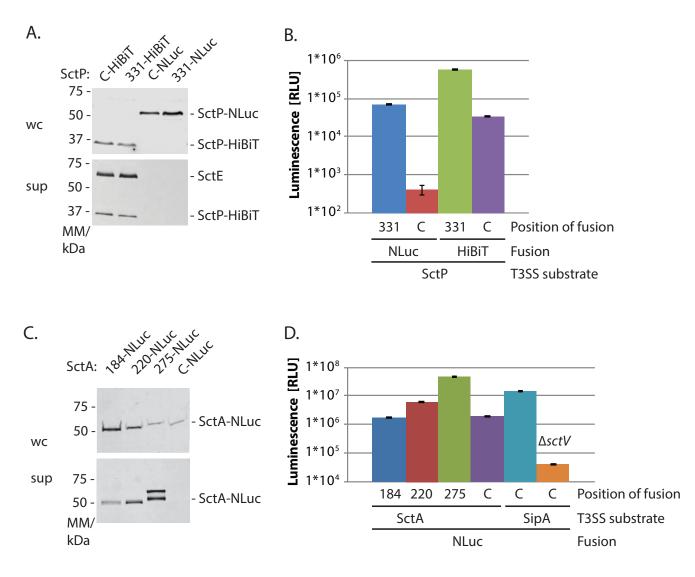
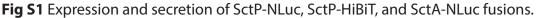


Figure 6





(A) Immunodetection of the indicated SctP-NLuc and SctP-HiBiT fusions, and of SctE on Western blot of SDS PAGE-separated culture supernatants and whole cell lysates. 331 means that NLuc or HiBiT was inserted behind residue 331 of SctP, so that the Shine-Dalgarno sequence of *sctQ*, which is overlapping with the gene of SctP, was unaffected. Note that SctE is not secreted when expressing SctP-NLuc fusions, i.e. SctP-NLuc cannot complement the needle length regulating function of SctP, thus substrate specificity switching to the secretion of intermediate substrates is not induced. (B) Luminescence of the indicated SctP-NLuc/HiBiT-fusions secreted into the culture supernatant. Data represent the mean (± standard deviation) of three technical replicates. Note that SctP₃₃₁-NLuc can be detected in the culture supernatant by luminometry but not by Western blotting. Also note that split-NLuc generally gives lower luminescence than regular NLuc.

(C) Immunodetection of the indicated SctA-NLuc fusions on Western blot of SDS PAGE-separated culture supernatants and whole cell lysates. The numbers (184, 220, 275) mean that NLuc was inserted behind these residues of SctA. The insertion positions where chosen based on the structure of *S*. Typhimurium SctA-1. Secreted SctA₂₇₅-NLuc reproducibly appeared as a double band for unknown reasons.

(D) Luminescence of the indicated SctA-NLuc and SipA-NLuc-fusions secreted into the culture supernatant. Data represent the mean (± standard deviation) of three technical replicates. Note that SctA_c-NLuc can be detected in the culture supernatant by luminometry but not by Western blotting. Also note that internal fusions of NLuc are acommodated well, with SctA₂₇₅-NLuc providing even stronger signal than SipA-NLuc.

Abbreviations: sup: culture supernatant, wc: whole cell lysates, C: C-terminus, RLU: relative luminescence units, NLuc: NanoLuc luciferase, T3SS: type III secretion system

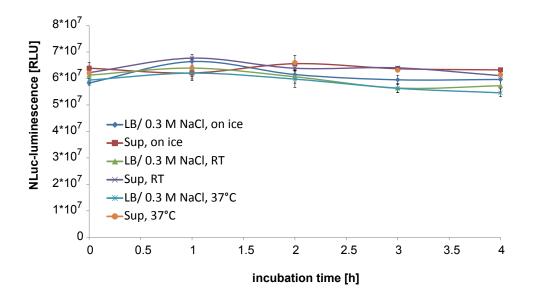


Fig S2 Stability of NLuc in LB/ 0.3 M NaCl and in culture supernatant. The enzymatic activity of purified NLuc was determined after incubation for 4 h at different conditions (on ice, room temperature (RT) and 37°C) in fresh LB/ 0.3 M NaCl and in filtered culture supernatant.