

1 A preliminary study of resting brain metabolism in
2 treatment-resistant depression before and after treatment
3 with olanzapine-fluoxetine combination

4 Short title: FDG PET in treatment-resistant depression with
5 olanzapine/fluoxetine combination therapy

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19 **Abstract**

20 Treatment-resistant depression (TRD) occurs in many patients and causes high
21 morbidity and mortality. Because TRD subjects are particularly difficult to study
22 especially longitudinally, biological data remain very limited. In a preliminary study to
23 judge feasibility and power, 25 TRD patients were referred from specialty psychiatric
24 practices. All were severely and chronically depressed and mostly had comorbid
25 psychiatric disorders as is typical in TRD. Nine patients were able to complete all
26 required components of the protocol that included diagnostic interview; rating scales;
27 clinical magnetic resonance imaging; medication washout; treatment with maximally
28 tolerated olanzapine-fluoxetine combination for 8 weeks; and pre- and post-treatment
29 fluorodeoxyglucose positron emission tomography. This drug combination is an
30 accepted standard of treatment for TRD. Dropouts arose from worsening depression,
31 insomnia, and anxiety. One patient remitted; three responded. A priori regions of
32 interest included the amygdala and subgenual cingulate cortex (sgACC; BA25).
33 Responders showed decreased metabolism with treatment in the right amygdala that
34 correlated with clinical response; no significant changes in BA25; better response to
35 treatment the higher the baseline BA25 metabolism; and decreased right ventromedial
36 prefrontal metabolism (VMPFC; broader than BA25) with treatment which did not
37 correlate with depression scores. The baseline metabolism of all individuals showed
38 heterogeneous patterns when compared to a normative metabolic database. Although
39 preliminary given the sample size, this study highlights several issues important for
40 future work: marked dropout rate in this study design; need for large sample size for
41 adequate power; baseline metabolic heterogeneity of TRD requiring careful subject

42 characterization for future studies of interventions; relationship of amygdala activity
43 decreases with response; and the relationship between baseline sgACC and VMPFC
44 activity with response. Successful treatment of TRD with olanzapine-fluoxetine
45 combination shows changes in cerebral metabolism similar to those seen in treatment-
46 responsive major depression.

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62 Introduction

63 Treatment-resistant depression (TRD) is often operationally defined as a major
64 depressive episode that fails to remit after treatment with at least two antidepressants of
65 different classes at therapeutic doses for an adequate treatment period [1-8]. More
66 extensive staging criteria for TRD have been proposed as well [1]. TRD must be
67 distinguished from inadequately treated depression resulting from numerous factors
68 such as patient non-compliance, intolerance to side effects, misdiagnosis (e.g., thyroid
69 disease), low dosage, etc. [2]. It is unclear if TRD is a particularly malignant form of
70 depression with its own pathophysiology, or if treatment-related changes in brain
71 metabolism are different than those found in treatment-responsive depression [3].
72 Patients with TRD are at increased risk of relapse [4]. Also, most TRD patients have
73 numerous psychiatric comorbidities inherently raising potential confounds with diagnosis
74 [5-8].

75 The STAR*D trial documents TRD is a frequent occurrence and a serious problem in
76 psychiatry afflicting about 30% of patients [4]. In terms of disease burden, it is second
77 only to back pain in terms of life-years of disability [9]. The high significance of TRD has
78 prompted an aggressive search for novel, more effective treatments including new
79 classes of antidepressants (e.g., glutamatergic receptor antagonists, neuroactive
80 steroids), add-on treatments (drug combinations, boosters such as lithium or
81 liothyronine, atypical neuroleptics, mood stabilizers), and devices for neuromodulation
82 (e.g., transcranial magnetic stimulation, vagus nerve stimulation, deep brain stimulation,
83 direct current stimulation).

84 Despite the clinical importance of TRD, few studies have examined its biology and
85 treatment. Several challenges have hindered such research. The recruitment of TRD
86 patients is difficult. These patients are likely heterogenous in pathology, have numerous
87 comorbidities, and are quite ill with significant risk for suicide. Their cross-sectional
88 physiology may be confounded by the many previous treatment trials. At best, the
89 medley of ineffective medications needs wash out, but this may lead to potential
90 withdrawal symptoms or symptom worsening. Encouragement to undergo yet another
91 treatment after so many failed trials becomes paramount. These patients need frequent
92 follow-up and clinician availability. Despite these impediments, some work has been
93 done using F¹⁸-fluorodeoxyglucose positron emission tomography (FDG PET) in those
94 with TRD [10-18]. However, not only does the biology of TRD, if homogeneous, remain
95 unclear, but also no biomarkers predicting treatment resistance have reached clinical
96 utility for this group of patients.

97 Past neuroimaging studies of treatment-responsive depression have highlighted the
98 amygdala and subgenual anterior cingulate cortex (sgACC) as key nodes of
99 depression-related circuitry showing reduction in activity with successful treatment (see
100 reviews [19-21]), although there are exceptions (e.g., no sg ACC change [22]; no
101 amygdala change [23]). Studies have suggested several features may characterize
102 TRD such as sgACC hyperactivity, amygdala hyperactivity, prefrontal/thalamic
103 dysconnectivity, habenular connectivity, prefrontal hypoactivity, and hippocampal
104 subfield volumes [10, 13, 14, 24-26]; however, consensus is yet to be achieved.

105 To our knowledge, there are no prior studies of brain metabolism in TRD patients
106 with drug washout to establish a baseline examined both before and after a full trial of a
107 combination antidepressant and atypical neuroleptic. Yet, the combination of
108 antidepressant and atypical neuroleptic is being used increasingly throughout the world
109 to treat TRD. The present study is of necessity preliminary, as no prior data existed with
110 which to power the sample size or even to determine its feasibility.

111 With a focus on the amygdala and sgACC based on prior literature, this cohort
112 observational study sought to test feasibility and to characterize regional brain
113 metabolism in TRD before and after adequate treatment with a combination of an
114 atypical neuroleptic (olanzapine) and fluoxetine (O/F), a selective serotonin reuptake
115 inhibitor (SSRI) antidepressant. Use of these drugs in combination will be referred to
116 hereafter as O/F.

117 This drug combination was the first drug approved by the USA Food and Drug
118 Administration in 2009 specifically for the indication of TRD [27]. There have been
119 several clinical trials, reviews, and meta-analyses examining the efficacy of O/F for
120 TRD; that discussion is beyond the scope of this study [28-35]. Originally, use of O/F
121 was based on preclinical work. Neurobiological changes associated with O/F dosing in
122 rats include increased prefrontal monoamine levels [36, 37] and suppression of limbic
123 immediate-early gene expression [38] relative to olanzapine or fluoxetine alone. No
124 differential effects on limbic neurogenesis were found using O/F, whereas the individual
125 drugs are associated with neurogenesis [39]. Whereas higher doses of O/F increased
126 levels of neurotrophin-3 selectively in rat prefrontal cortex, low O/F doses or higher

127 doses of olanzapine or fluoxetine administered individually did not [40]. These animal
128 findings suggest some unique effects of O/F therapy not accounted for by the actions of
129 the individual drugs.

130 The aim of the present work was not to assess efficacy or health outcomes in a
131 clinical trial, as this has been reported previously (see above). Rather, the project's
132 purpose was to use imaging to bear on the question of mechanisms. Whether such
133 combination treatments for TRD follow similar metabolic effects to other antidepressants
134 in treatment-responsive depression is unknown. In addition, the individual TRD patient's
135 deviation from a normative database indicate for the first time the potential changes in
136 regional brain metabolism in an individual, unmedicated, TRD patient. Such data could
137 address preliminarily whether sgACC hypermetabolism or other biomarker is
138 characteristic of TRD, a key issue in patient selection for future treatment trials of TRD.

139 **Materials and Methods**

140 **Participants**

141 Twenty-five participants with severe TRD were enrolled. They had many
142 psychotherapy and medication trials, some even failing convulsive therapy. They all had
143 a longstanding chronic illness lasting many years. Most had comorbid psychiatric
144 disorders. They were recruited and enrolled through referral from physicians' outpatient
145 clinics known to specialize in the treatment of TRD (co-authors: DA, BR, FSA). The
146 principal inclusion criterion was severe, refractory major unipolar depressive disorder
147 (Scheduled Clinical Interview for Diagnostic and Statistical Manual-IV; SCID [41]) as the

148 primary diagnosis. Exclusion criteria included a lifetime history of cognitive impairment,
149 psychosis, bipolar disorder, drug dependence, pregnancy, as well as any clinically
150 significant findings on magnetic resonance imaging. All subjects provided written
151 informed consent as approved by the VA Institutional Review Board (IRB) and the
152 Radioactive Drug Research Committee (RDRC) approved by the FDA.

153 **Treatment**

154 Patients' polypharmacy was tapered with washout for two weeks before the baseline
155 measurement of glucose uptake using FDG PET as described previously [42]. They
156 were then titrated to the maximal tolerated dose of fluoxetine (≤ 60 mg) and olanzapine
157 (≤ 30 mg). The use of the combination of an SSRI such as fluoxetine and an atypical
158 neuroleptic such as olanzapine is a standard of care in the management of patients with
159 TRD. The maximal tolerated dose was held constant for eight weeks except for rescue
160 medication of low dose benzodiazepine or short acting hypnotics for severe anxiety or
161 insomnia. Given the seriousness of the illness, risk for suicide, and focus on
162 mechanisms rather than efficacy, the study had no placebo control medication. Patients
163 were seen either weekly or every two weeks (depending on stability) during wash out
164 and treatment. Compliance was checked by counting of pills every 1-2 weeks. Side
165 effects were evaluated with open-ended questions without checklists typical of clinical
166 trials. After treatment with O/F for eight weeks and PET imaging, they were returned to
167 their referring psychiatrist for continued assessment and follow-up.

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169 **Clinical Assessments**

170 All subjects were assessed by their referring physician as having TRD as their
171 primary diagnosis. Medical records were reviewed to ensure all subjects had at least
172 two trials of antidepressants from different classes with adequate doses and duration of
173 treatment (at least stage III TRD [1]). All subjects underwent structured diagnostic
174 interviews using the SCID-1, Clinician Version [43]. The primary outcome measure was
175 the change in Montgomery-Asberg Depression Rating Scale (MADRS [44]) score after
176 eight weeks of treatment. A clinical response was defined as a greater than or equal to
177 50% drop in depression score, while a remission was defined as a MADRS of less than
178 or equal to 8. Anxiety was scored with the Hamilton Anxiety Scale (HAMA [45]).
179 Additional testing not directly pertinent to the present study included Clinical Global
180 Impression Scale (CGI [46]), Mini-Mental Status Exam (MMSE [47]), Shipley Institute of
181 Living Scale [48], Edinburgh Handedness Inventory [49], Profile of Mood States (POMS)
182 [50], and Positive Affect Negative Affect Scale (PANAS [51]).

183 **Positron emission tomography**

184 Patients were scanned after washout (pre-treatment or baseline) and after
185 completion of the eight weeks of treatment at maximal tolerated dose (post-treatment).
186 Participants fasted for at least six hours before imaging; blood glucose was checked
187 immediately before scanning. The relative regional glucose uptake was measured by
188 injecting intravenously a bolus of ^{18}F -FDG in saline at a dose of 185 MBq (5 mCi)/70 kg.
189 They rested for 50 minutes during tracer uptake with eyes closed and ears open in a

190 dimly lit, quiet room while being monitoring for wakefulness. The scanner was a
191 Siemens (Knoxville, TN, USA) ECAT EXACT 47 operated in 2-D mode with septae
192 extended. After measured attenuation, counts were collected during an emission scan
193 lasting 20 minutes. Data were corrected for scatter, decay, randoms, and electronic
194 deadtime. Images were reconstructed using filtered backprojection to a resolution of
195 approximately 12 mm full width at half maximum.

196 **Image analysis**

197 Image analysis followed routine procedures including normalization for whole-brain
198 activity, intersubject stereotactic averaging, subtraction of pre- from post-treatment
199 activity, statistical parametric mapping, and threshold $Z = 3.3$ as previously described
200 [52]. Images were warped nonlinearly into stereotactic space [53] with regression for
201 age using Neurostat software [54]. Parametric maps were overlaid on a template MRI
202 blurred to a similar resolution as the PET scan. In-house software (iiV,
203 <http://james.psych.umn.edu/iiV/>) was used to display results on a standard MRI
204 template [55]. For an exploratory look at individual TRD patient's whole-brain voxelwise
205 differences from our normative database ($N = 30$). For the purposes of display, the
206 individually warped t images used an uncorrected magnitude threshold of $p < \sim 0.05$.

207 **Regions of interest**

208 Two ROIs were examined based upon existing, extensive literature (see Fig 1):
209 amygdala and sgACC. The amygdala ROI consisted of a sphere of 13 mm diameter
210 center on each amygdala in the atlas of Talairach and Tournoux (1988, Fig 1) at

211 coordinates (± 23 , -4, -16). Mean counts were collected for each ROI from the pre- and
212 post- treatment scans for each patient. The percent change in MADRS score was
213 regressed linearly against the mean amygdala activity. Paired t-tests were used to
214 compare mean amygdala counts before and after treatment for the group and for
215 responders vs. non-responders separately. A threshold of $p < 0.05$ was used for the
216 ROI analysis.

217 

218 **Figure 1. Amygdala (A) and sgACC/VMPFC (B) regions of interest.**

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220 Based upon an *a priori* focus on the sgACC, we defined two additional subregions
221 within sgACC for this analysis. Previous studies point to changes in subgenual
222 metabolism that are not restricted to the sgACC of Brodmann area 25 (BA25) but
223 extend along the ventromedial cingulate cortex [10, 18, 56, 57]. So, a region including
224 only those voxels labeled as BA25 by the atlas of Talairach and Tournoux [53] was
225 created using the Talairach Daemon
226 (<http://ric.uthscsa.edu/projects/talairachdaemon.html>) [58]. A second, less specific
227 region was drawn as a cuboid on each hemisphere with the following extents measured
228 from the anterior commissure: x, $\pm(1-12$ mm); y, 10-25 mm; z, -5 to -16) mm [53]. This
229 approximated the region of hypometabolism associated with antidepressant treatment
230 reported previously by our laboratory [18]. The region encompasses BA25, as well as
231 portions of BA32 and BA33 and will be referred to as subgenual/VMPFC (Fig 1B).

232 Mean counts from each of these subgenual regions were compared pre- and post-
233 treatment for responders and non-responders using paired *t*-tests. Percent-change in
234 mean ROI metabolism was linearly regressed against percent-change in MADRS
235 across all subjects. Pre-treatment mean metabolism was also regressed against
236 percent change in MADRS.

237 One additional unplanned ROI was included post hoc because of its proximity to the
238 amygdala, involvement in depression, and similar response to treatment in published
239 work [22]. A 13 mm diameter spherical ROI was centered on the hippocampus at the
240 following Talairach coordinates: x, ± 27 mm, y, -23 mm; z, -9 mm or MNI (± 27 , -22, -0)
241 <http://sprout022.sprout.yale.edu/mni2tal/mni2tal.html>.

242 A repeated measures ANOVA was performed on the amygdala, VMPFC, and head
243 of the hippocampus ROIs as defined above. Extensive prior literature highlights these
244 regions as important in affective illness. The BA25 region was not included as no
245 change in activity occurred with treatment (see below). The dependent measure was
246 glucose uptake (PET counts). Repeated measures included TIME (Pre-treatment, Post-
247 treatment), ROI (amygdala, hippocampus, and VMPFC), and SIDE (right hemisphere,
248 left hemisphere). Additionally, the correlation matrix for glucose uptake across
249 amygdala, VMPFC, and hippocampus was calculated (S2 Table).

250 **Results**

251 **Clinical response**

252 Drug wash out led frequently to increased insomnia, anxiety, and depression which were
253 the leading reasons for discontinuation by participants (N = 16). Side effects from O/F observed
254 reflected those commonly seen with this drug combination [28]. There were no suicide attempts
255 during the study. Nine participants completed the entire protocol (clinical MRI; drug washout;
256 O/F treatment; Pre- and Post-treatment PET scans). Clinical data are shown in Table 1 which
257 includes demographics, gender, family history of depression, illness onset, comorbidities (both
258 current and lifetime), failed treatments, and response designation. The HAMA and MADRS
259 scores as well as weights before and after treatment are displayed in S1 Table. The average
260 tolerated dose was 39 mg (range 30-45 mg) of fluoxetine and 12 mg (range 10-12 mg) of
261 olanzapine. Their weight increased significantly during treatment (S1 Table; S1 Figure); there
262 was no interaction between time (pre vs post) and response ($p > 0.2$). The average baseline
263 MADRS score was 31 (SD 5; range 24-38); no significant differences arose in baseline MADRS
264 score between responders vs. non-responders ($p > 0.3$; no shown). Completers showed a
265 decrease in MADRS score following eight weeks of treatment (Mean, 12; SD, 5; S1 Table). The
266 baseline HAMA did not differ between responders vs. non-responders ($p < 0.72$). The HAMA
267 declined significantly also after treatment ($t(8) = 4.7$; $p < 0.002$) with the significance driven by
268 the non-responders ($p < 0.006$; responders, $p < 0.13$). Four patients responded; one of these
269 achieved remission. Of note, responders tended to have fewer comorbidities and earlier onset
270 than non-responders.

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276 **Table 1. Patient demographics and outcomes.**

Age	Sex	Family History of MDD	Life time Comorbidity	Current Comorbidity	Failed medications	Baseline MADRS	% Change	Response
47	F	+		OCD, Anorexia, PD, Anxiety NOS, Dysthymia	SSRIs, venlafaxine, atypical neuroleptics, TCAs, mirtazapine, nefazodone, bupropion, ECT, anticonvulsants, buspirone	33	-6%	-
44	M	+		PD, Dysthymia	SSRIs, venlafaxine, mirtazapine, benzodiazepines	29	-17%	-
52	M	unknown			SSRIs, venlafaxine, lithium	37	-22%	-
36	M	+	OCD, ADD		SSRIs, venlafaxine, TCA, MAOI, T3, stimulants, dopamine agonist, anticonvulsants	27	-26%	-
62	F	+			SSRIs, venlafaxine, anticonvulsant, atypical neuroleptic	29	-59%	+
54	M	+			SSRIs, venlafaxine, lithium, buspirone, TCA, anticonvulsants	24	-83%	+(remitted)
61	M	+	OCD, ADD, dysthymia		SSRIs, venlafaxine, bupropion, benzodiazepine, stimulants, T4, nefazodone	31	-35%	-
27	M	+	ADHD	GAD	SSRIs, venlafaxine, atypical neuroleptic, mirtazapine	38	-50%	+
41	M	+			SSRIs, venlafaxine, atypical neuroleptic	28	-64%	+

277 M, male; F, female; MDD, major depressive disorder, OCD, obsessive compulsive disorder; ADD,

278 attention deficit disorder; ADHD, attention deficit disorder with hyperactivity; PD, personality disorder;

279 NOS, not otherwise specified; GAD, generalized anxiety disorder; SSRI, selective serotonin reuptake
280 inhibitor; TCA, tricyclic antidepressant; ECT, electro-convulsive therapy; MAOI, monoamine oxidase
281 inhibitor; T3, liothyronine; T4, levothyroxine; MADRS, Montgomery Asberg Depression Rating Scale

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283 A 2 x 2 Analysis-of-Variance with Treatment-response (responder vs. non-
284 responder) as the between-subject factor and Time (post vs. pre-test) as the within-
285 subject factor demonstrated a main effect of Time ($F(1,7) = 195.2, p < 0.001$), confirming
286 that MADRS scores decreased significantly across sessions. The interaction of
287 Treatment-response X Time was also significant ($F(1,7) = 43.9, p < 0.001$). This
288 interaction confirmed that responders showed significantly greater reductions in MADRS
289 scores than non-responders.

290 Whole-brain exploratory image analysis

291 Voxel-wise comparisons of patients' post- and pre-treatment scans revealed
292 significant changes (Table 2; S2-4 Figures). Patients were grouped as responders and
293 non-responders (see above) to identify metabolic change associated with successful
294 treatment.

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Table 2. Significant changes in whole-brain voxelwise analysis.

X (mm)	Y (mm)	Z (mm)	Z-score	Region
<i>All patients</i>				
12	-35	54	+3.4	R paracentral lobule
6	-60	43	+3.3	R precuneus (BA7)

-12	-60	-20	-4.6	L Cerebellum
17	-94	-4	-3.7	R Lingual Gyrus (BA18)
28	14	-20	-3.6	R Inferior Frontal Gyrus (BA47)
-15	-33	7	-3.6	L Thalamus
-37	-64	-22	-3.5	L Cerebellum
30	-49	-27	-3.4	R Cerebellum
-12	-10	0	-3.4	L Thalamus
10	-1	-4	-3.3	R globus pallidus
Non-responders				
-8	-60	-16	-5.2	L Cerebellum
8	-49	-29	-4.6	R Cerebellum
-17	-53	29	-4.4	L Cingulate Gyrus (BA31)
-44	32	7	-3.9	L Inferior Frontal Gyrus (BA45)
-10	-10	2	-3.9	L Thalamus
17	-91	-2	-3.7	R Lingual Gyrus (BA17)
-51	-67	14	-3.7	L Middle Temporal Gyrus (BA39)
-21	-73	-18	-3.7	L Cerebellum
1	-19	-16	-3.4	R Midbrain
-42	-4	38	-3.4	L Precentral Gyrus (BA6)
Responders				
-48	-13	2	+3.5	Left Superior Temporal Gyrus (BA22)
-53	-24	18	+3.4	Left Post-Central Gyrus (BA40)
6	-49	36	+3.3	Right Precuneus (BA31)
10	-58	38	+3.3	Right Precuneus (BA7)
19	-1	-7	-4.4	Right Dorsal Amygdala

R, right; L, left; BA, Brodmann area

299 Responders showed increases in the left superior temporal and post-central gyri and
300 right precuneus. Significantly reduced metabolism was confined to a peak in the right
301 amygdaloid complex (see Fig 2). Of note, liberalizing the threshold to $0 < 0.05$
302 (uncorrected) revealed bilateral amygdala deactivations. Non-responders showed no
303 significant increases in metabolism following treatment. Areas of reduced metabolism
304 after treatment were found in non-responders in the bilateral cerebellum, lingual gyrus,
305 middle temporal gyrus, thalamus, midbrain, and dorsal cingulate gyrus (see Table 2).
306 For all patients as a group, no significant changes occurred in limbic structures in the
307 whole-brain image analysis.

308

309 **Figure 2. Right amygdala metabolism in responders following**
310 **olanzapine/fluoxetine treatment was the only significant decrease in the whole-**
311 **brain voxelwise analysis.** Upper left section, coronal; lower section, transverse; upper
312 right section, sagittal.

313

314 No studies have reported on individual metabolic patterns associated with
315 unmedicated TRD. To explore this variability while accepting the limitation of the risk for
316 false positive or negative responses, each TRD subject's baseline warped FDG PET
317 (i.e., after medication washout or baseline) was contrasted with those of a normative
318 database with threshold set at $t=2.0$ for visualization as performed previously [42].
319 These individual subtractions for the nine completers are show in S5 Figure including
320 both increases and decreases in metabolism. Examination of individual subject's scans

321 showed considerable heterogeneity in baseline scans with a mixture of positive,
322 negative, or null changes in the BA25/VMPFC region. This variability suggests baseline
323 heterogeneity in TRD or metabolic changes related to the previous history of failed
324 treatments for TRD. If any antidepressant response occurred, a relationship to
325 metabolic signature was not evident.

326 **Region of interest analyses**

327 **ANOVA on amygdala, hippocampus, and VMPFC**

328 The repeated measures ANOVA indicated significant main effects of TIME ($F(1,8) =$
329 $5.992, p = 0.04$) and ROI ($F(2,16) = 24.94, p = 0.001$) without significant interaction
330 effects (S6 Figure). This result indicates that O/F reduces activity in all three ROIs. The
331 analysis of the correlation matrix for the ROIs reached significance only for the
332 correlation between the left and right sgACC post-treatment ($r = 0.82, p = 0.007$; S2
333 Table). O/F tended to increase each region's inter-hemispheric functional connectivity
334 compared to baseline.

335 **Amygdala region**

336 Responders showed a significant reduction in mean glucose metabolism in the right
337 amygdala ROI ($t(3) = 3.38, p = 0.04$), while non-responders showed no change with
338 treatment ($t(4) = -0.68, p = 0.54$) (Fig 3A). There was no change in left amygdala
339 metabolism in either group (responders: $t(3) = 0.88, p = 0.45$; non-responders: $t(4) =$
340 $1.29, p = 0.28$) (Figure 3A). Regression analyses showed no correlation between
341 percent reduction in left amygdala metabolism and percent-reduction in MADRS scores

342 across all subjects (Fig 3B, $r = -0.20$, $p = 0.61$). This correlation was significant in the
343 right amygdala (Fig 3B, $r = -0.70$, $p = 0.03$). Because the hippocampus is near the
344 amygdala and the resolution of PET in this study is low, the post hoc hippocampal
345 region was also examined for changes in activity.

346

347 **Figure 3. Changes in amygdala glucose uptake before and after treatment and its**
348 **relationship to depression symptoms.** (A) Amygdala metabolism examined
349 separately for responders vs. non-responders in the right and left amygdala before and
350 after treatment. ^a $p < 0.03$. (B) Change in MADRS scores and change in amygdala
351 metabolism. Only the right amygdala showed a significant correlation between change
352 in metabolism and change in MADRS scores. Red line identifies threshold for response.

353

354 **Exploratory subgenual regions (BA 25 & sgACC/VMPFC)**

355 Paired t -tests revealed no changes with treatment in either the left or right BA25
356 metabolism in either responders (left: $t(3) = 0.05$, $p = 0.97$; right: $t(3) = 0.71$, $p = 0.53$) or
357 non-responders (left: $t(4) = 1.13$, $p = 0.32$; right: $t(4) = 0.67$, $p = 0.54$). Changes in left or
358 right BA25 metabolism showed no correlation with changes from baseline MADRS (left:
359 $r = 0.29$, $p = 0.45$; right: $r = 0.15$, $p = 0.69$). However, baseline right hemisphere BA25
360 metabolism showed a marginally significant correlation with change from baseline
361 MADRS, whereby higher baseline metabolism predicted better response to O/F therapy

362 ($r = 0.68$, $p = 0.05$). This correlation was not significant in the left hemisphere ($r = 0.13$,
363 $p = 0.74$).

364 Paired t-tests revealed a significant decrease in right but not left VMPFC metabolism
365 in responders (left: $t(3) = 0.07$, $p = 0.95$; right: $t(3) = 4.18$, $p = 0.02$), and no significant
366 change in VMPFC metabolism in non-responders (left: $t(4) = 0.81$, $p = 0.47$; right: $t(4) =$
367 0.53 , $p = 0.62$). Changes in VMPFC metabolism showed no correlation with change in
368 MADRS (left: $r = 0.11$; $p = 0.77$; right: $r = 0.18$, $p = 0.64$). Baseline metabolism in the
369 right, but not left, VMPFC area correlated with MADRS reduction, with higher baseline
370 metabolism predicting better response (left: $r = 0.37$, $p = 0.33$; right: $r = 0.84$, $p < 0.01$).

371 Discussion

372 This preliminary study found that medication-free TRD patients treated with O/F at
373 therapeutic doses for an adequate duration showed a response-related decline in the
374 metabolism of the right dorsal amygdala using a whole-brain, voxel-wise analysis. The
375 dorsal amygdala in humans consists mostly of the central nucleus, the terminus of the
376 spino-parabrachial-amygdaloid pain pathway and the principal efferent pathway for the
377 emotional and physiological processing. Given the low resolution of the present study,
378 caution is warranted in localization pending higher resolution techniques (e.g., higher
379 resolution PET scanners and coregistered high resolution MRI). Several increases in
380 metabolism surfaced also after treatment.

381 Two regions of interest (and one post-hoc region) based on existing literature were
382 examined for drug-related changes in metabolism and relationship to response.

383 Following treatment, the change in metabolism of the right amygdala correlated with the
384 change in MADRS score; no correlation was observed for the left amygdala. Treatment
385 did not significantly change the smaller sgACC region defined as BA25. However, a
386 broader ROI along the VMPFC approached significance despite the small sample size.

387 ANOVA of ROIs in the VMPFC, amygdala, and immediately adjacent hippocampus
388 (a post hoc exploratory ROI) indicated a main effect of treatment and ROI without
389 significant interactions. The VMPFC showed the highest activity. O/F was associated
390 with reduced activity in all ROIs tested. The hippocampus did not appear responsible for
391 the deactivation seen in the dorsal amygdala. However, the hippocampus may have
392 followed similar changes with treatment that did not reach significance; such changes
393 have been reported previously.

394 Baseline metabolism in TRD may have relevance to treatment response and may, if
395 confirmed, find utility for patient selection in treatment trials. Greater baseline
396 metabolism in right BA25 and sgACC/VMPFC predicted better response to O/F therapy.
397 Also, responders showed a significant decrease in metabolism following treatment in
398 the right VMPC region which could reflect a higher initial baseline (as BA25 was
399 included in the right VMPFC region). The higher baseline BA25 metabolism in
400 responders to O/F may relate to 1) higher baseline resting sgACC/VMFPC blood flow
401 seen in TRD patients when compared to controls during neuromodulation trials [10, 11];
402 2) increased sgACC blood flow induced by sadness induction [59]; and 3) and
403 increased resting blood flow in sgACC/VMFPC in healthy subjects high in negative
404 affect [60]. Likewise, the decline in right VMPFC activity in TRD treated with O/F is

405 analogous to the decline in resting blood flow in VMPFC in TRD responders treated with
406 deep brain stimulation [10] as well as decreased VMPFC metabolism with chronic
407 vagus nerve stimulation [18, 56].

408 Strengths of the study include the recruitment of severely ill TRD patients,
409 medication washout before treatment, exploratory examination of individual TRD
410 patients, and full course of treatment with O/F to maximal tolerated doses. These
411 preliminary data suggest heterogeneity in metabolic signatures in TRD. If replicated, this
412 heterogeneity requires consideration in the design of trials for TRD using medications or
413 devices. However, the changes observed with treatment appear broadly like those
414 reported in treatment-responsive major depression with other antidepressant
415 treatments: decreased amygdala and VMPFC metabolism.

416 Limitations of this study include comorbidities, prior heterogeneous treatments
417 during past failed trials, significant participant dropout, small final sample size, lack of a
418 placebo, low PET resolution, and confounding of depression with anxiety measures.
419 Patient dropouts and comorbidities could limit generalizability. However, most TRD
420 patients do have comorbidity [5-8]. A larger replication sample could address whether
421 major comorbidities represent a covariate of interest in the response as suggested by
422 this study—likewise for dropouts. Future studies will need to account for the large
423 dropout during washout. The limited resolution of PET places some ambiguity in the
424 precise determination of amygdala activity. However, the latest scanners with resolution
425 of 3 mm full width at half maximum will improve localization in future studies. Of note,
426 although the ROI analysis of the right amygdala confirmed significant deactivation after

427 treatment that correlated with clinical response, the focus was only partially resolved
428 from other nearby regions which showed a similar response pattern (e.g.,
429 hippocampus). Depression and anxiety are both prominent symptoms in TRD, and
430 antidepressants/atypical neuroleptics often decrease both depression and anxiety
431 scores. In this regard, HAMD and HAMA scores are inter-correlated [61]. Anxiety scores
432 measured in this study in responders before and after treatments did not differ.
433 Therefore, the decrease in amygdala activity does not likely reflect changes in anxiety,
434 but rather depression. The severity of illness and risk of suicide precluded a placebo
435 control. However, other studies using FDG PET and placebos in test-retest designs
436 suggest high consistency in normalized regional activity and only small changes [62-64].

437 **Conclusions**

438 TRD patients show considerable baseline metabolic heterogeneity following
439 medication washout. Whether this heterogeneity arises from differing disease
440 pathologies or from effects of past treatments remains unclear. As reported for other
441 antidepressant therapies in treatment-responsive depression, decline in amygdala and
442 VMPFC activity surfaced here with O/F treatment. Furthermore, decreased metabolism
443 in the right amygdala with treatment correlated with improvement in depression
444 following O/F treatment.

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455 **Conflicts of Interest**

456 The authors declare no conflict of interest.

457 **Authors' Contributions**

458 JVP designed the study, secured funding, participated in data collection
459 and analyses, and wrote/edited the final manuscript. SAS designed the
460 study, assisted in securing funding, participated in data collection and
461 reviewed and edited the final manuscript. GS analyzed data, provided
462 figures, wrote the initial manuscript, and reviewed and edited the final
463 manuscript. JTL analyzed data, contributed software, curated data,

464 provided figures, and edited the final manuscript. DA, BR, FSA provided
465 clinical care and recruitment of the patients, gathered clinical data, and
466 edited the final version of the manuscript.

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657 **Supporting Information**

658 **S1 Figure. Weights before and after O/F combination.**

659 **S2 Figure. Brain glucose uptake in non-responders: Post- minus Pre- O/F**
660 **treatment.**

661 Stereotactically normalized. Image left is right side of brain. AC-PC plane 0 mm. Color
662 scale shows Z-scores with threshold $Z = \pm 3.3$.

663 **S3 Figure. Brain glucose uptake in responders: Post- minus Pre- O/F treatment**

664 Stereotactically normalized. Image left is right side of brain. AC-PC plane 0 mm. Color
665 scale shows Z-scores with threshold at $Z = \pm 3.3$.

666 **S4 Figure. Brain glucose uptake for all subjects: Post- minus Pre- O/F treatment.**

667 Stereotactically normalized. Image left is right side of brain. AC-PC plane 0 mm. Color
668 scale shows Z-scores with threshold at $Z = \pm 3.3$.

669 **S5 Figure. Differences in resting brain glucose uptake between individual**
670 **subjects (N = 9) at baseline (after washout) and a normative data set (N = 30).**

671 For visualizing individual metabolic fingerprints of all nine subjects, the threshold was
672 set at $t = 2.0$ that is the usual threshold used for studying change in individuals [42].

673 Each subject is represented by a study number (e.g., pL0009). Age regression was
674 used to match individual subject's age to that of the normative group. R, right; L, left, A,
675 anterior; P, posterior. The patterns are heterogenous. For example, some individuals
676 have sgACC/VMPFC hypoactive, hyperactivity, or no change.

677 **S6 Figure. Main effects of TIME and ROI on glucose uptake.**

678 **S1 Table. Individual subject's weight, depression scores, and anxiety ratings.**

679 **S2 Table. Correlation matrix for metabolism in bilateral ROIs.**

680 Green cells below diagonal are for Pre-treatment; blue cells above diagonal are for

681 Post-treatment. R, right; L, left; Hippo, hippocampus; sgACC, subgenual anterior

682 cingulate/VMPFC. † $p < 0.007$

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