

1 **Abundance of Nef and p-Tau217 in brains of individuals diagnosed with HIV-associated**
2 **neurocognitive disorders correlate with disease severance**

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15 Short title: p-Tau217 and lipid rafts accumulate in HAND brains

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20

21 **Abstract**

22 HIV-associated neurological disorders (HAND) is a term used to describe a variety of
23 neurological impairments observed in HIV-infected individuals. The pathogenic mechanisms of
24 HAND and its connection to HIV infection remain unknown. The brain samples from HIV-
25 infected individuals, both with and without HAND, were characterized by increased abundance
26 of p-Tau217 peptide, which correlated with the abundance of flotillin 1, a marker of lipid rafts.
27 HIV-1 Nef was detected in some, but not all, samples from HAND-affected individuals. Samples
28 positive for Nef had lower abundance of cholesterol transporter ABCA1, higher abundance of
29 flotillin 1 and p-Tau217, and were obtained from individuals with higher severity of HAND
30 relative to Nef-negative samples. These results highlight the contribution of Nef and Nef-
31 dependent effects on cholesterol metabolism and lipid rafts to the pathogenesis of HAND and
32 support a connection between pathogenesis of HAND and Alzheimer's disease.

33

34 Key words: HIV-1; HAND; Nef; p-Tau217; lipid rafts; ABCA1; cholesterol

35

36 **Introduction**

37 HIV-associated neurological disorders (HAND) have clinical hallmarks of a
38 neurodegenerative disease with progressive cognitive decline reflected initially by neuronal
39 dendritic simplification and ultimately by the neuronal loss [1]. With the introduction of
40 combination anti-retroviral therapy (cART), prevalence of the severe form of HAND, HIV-
41 associated dementia (HAD), has diminished, but that of milder forms, mild neurocognitive
42 disorder (MND) and especially asymptomatic neurocognitive impairment (ANI), have increased
43 [2], suggesting that cART slows progression, but does not prevent initiation of the disease. The
44 reasons for this persistence are not fully understood and several possible explanations have been
45 suggested (reviewed in [3, 4]). One of the mechanisms gaining support in the recent studies links
46 HAND pathogenesis to beta amyloid, suggesting a connection between HAND and Alzheimer's
47 disease (reviewed in [5]). HAND has many similarities to Alzheimer's disease (AD), such as
48 neuroinflammation, similar transcriptional signatures, and increased abundance and changed
49 localization of intracellular beta amyloid (A β), a critical component in AD pathogenesis [5-8].
50 While the possibility that HIV infection "speeds up" the underlying AD is an attractive
51 hypothesis, the mechanisms of this connection remain unclear. Described below are results of
52 our previous studies of the effects of Nef on cellular cholesterol metabolism and lipid rafts,
53 which provide a plausible mechanism.

54 Expression of Nef in HIV-infected cells induces degradation of the key cellular
55 cholesterol transporter ABCA1, causing suppression of cholesterol efflux and increasing
56 abundance of the lipid rafts [9, 10]. Most importantly, the same activity was demonstrated for the
57 Nef-carrying extracellular vesicles [11, 12], which continue to be released into circulation and
58 the brain from HIV-infected cells even after suppression of HIV replication by cART [7, 13, 14].

59 Amyloidogenic proteins concentrate in the lipid rafts [15], and most amyloid proteins involved in
60 pathogenesis of neurodegeneration are raft proteins [16-18]. High local concentration of
61 amyloidogenic proteins is a prerequisite for the effective nucleation and cascading progression of
62 their misfolding, a key pathogenic element of neurodegeneration. Therefore, upregulation of the
63 lipid rafts by Nef or Nef-containing extracellular vesicles and accumulation of amyloidogenic
64 proteins in the rafts may accelerate the progress of their misfolding and promote its spread
65 through the brain. A similar mechanism has been shown for prions, which also promote lipid raft
66 formation and depend on lipid rafts for progression of prion misfolding [19].

67 In this study, we demonstrated associations between the presence of Nef with reduced
68 levels of ABCA1, increased abundance of flotillin 1, disease severity, and the increased
69 abundance of p-Tau217 - which is a characteristic marker of the Alzheimer's disease [20-22].
70 Our findings support the proposed mechanism of HAND whereby Nef-mediated suppression of
71 ABCA1 increases the abundance of lipid rafts, which in turn enables the progression of
72 tauopathy.

73

74 **Materials and Methods**

75 *Brain samples:* Fresh-frozen post-mortem samples from hippocampus or mid-temporal gyrus of
76 HIV-infected individuals with or without HAND diagnosis were obtained from National
77 NeuroAIDS Tissue Consortium (NNTC). Brain samples from uninfected individuals without
78 neurodegenerative disease diagnosis were obtained from the National Institutes of Health
79 NeuroBioBank. All samples were deidentified, and provided clinical information is presented in
80 Table S1. The studies abided by the Declaration of Helsinki principles.

81

82 *Western blot analysis:* Samples were analyzed by automated Western immunoblotting using the
83 JessTM Simple Western system (ProteinSimple, San Jose, CA). For analysis of ABCA1, a 66-440
84 kDa Jess separation module (SM-W008) was used, for Nef – a 2-40 kDa module (SM-W012),
85 for p-Tau217 and Flotillin 1 – 12 - 230 kDa module (SM-W004). Three μ L of brain lysates (1
86 μ g/ μ L protein) was mixed with 1 μ L of Fluorescent 5X Master mix (ProteinSimple) in the
87 presence of fluorescent molecular weight markers and 400 mM dithiothreitol (ProteinSimple).
88 This preparation was denatured at 95°C for 5 min for Nef, p-Tau217 and Flotillin 1 analysis. For
89 ABCA1, the mix was incubated at 37°C for 15 min. Molecular weight ladder and proteins were
90 separated in capillaries through a separation matrix at 375 volts. A ProteinSimple proprietary
91 photoactivated capture chemistry was used to immobilize separated viral proteins on the
92 capillaries. Capillaries with immobilized proteins were blocked with KPL Detection Block (5X)
93 (SeraCare Life Sciences, Gaithersburg, MD, cat. #5920-0004) for 60 min, and then incubated
94 with antibodies for 60 min. After a wash step, HRP-conjugated goat anti-rabbit (cat. #042-206),
95 donkey anti-goat (cat. #043-522-2), or goat anti-mouse antibodies (cat. #042-205) from
96 ProteinSimple were added for 30 min. The chemiluminescent revelation was established with
97 peroxide/luminol-S (ProteinSimple). Digital image of chemiluminescence of the capillary was
98 captured with Compass Simple Western software (version 5.1.0, Protein Simple) that calculated
99 automatically heights (chemiluminescence intensity), area, and signal/noise ratio. Results were
100 visualized as electropherograms representing peak of chemiluminescence intensity and as lane
101 view from signal of chemiluminescence detected in the capillary. A total protein assay using
102 Total Protein detection module DM-TP01 and Replex Module RP-001 was included in each run
103 to quantitate loading.

104

105 *Antibodies:* For Nef detection, the following mouse monoclonal antibodies were used: 3D12
106 (ABCAM ab42355) against amino acids ³⁵RDLEKHGAISSNTAA⁵⁰ of HIV-1 HXB2; EH1
107 (AIDS reagents #ARP-3689) against ¹⁹⁴MARELHPEYYKDC²⁰⁶ of HIV-1 B subtype consensus;
108 SN20 (gift from Dr. Bernhard Maier, Indiana University) against the SH3 binding domain of Nef
109 (FPVTPQ); and 6JR (ABCAM ab42358) mapped to ¹⁹⁵ARELHPEYYKD²⁰⁵ of the HIV-1 B
110 subtype consensus. P-Tau217 was detected using the rabbit polyclonal antibody to Tau
111 phosphorylated on Threonine 217 (ThermoFisher (Waltham, MA), cat. #44744); ABCA1 – using
112 mouse monoclonal anti-ABCA1 antibody H10 (Abcam (Cambridge, MA), cat. #ab18180); and
113 flotillin 1 – using goat anti-flotillin 1 polyclonal antibody (Novus Biologicals (Littleton, Co), cat.
114 #NB1001043).

115

116 *Statistics:* Statistical analyses including Spearman correlations, Wilcoxon rank-sum tests,
117 Kruskal-Wallis test with DSCF multiple comparisons option, and multivariate linear regression
118 were conducted in SAS v9.4. Data visualization was conducted in GraphPad PRISM v.9. For
119 multivariate analysis, Box Cox transformation of variables was performed in SAS v9.4 prior to
120 regression. Figure captions describe the statistical test used.

121

122 **Results**

123 The role of amyloid proteins in pathogenesis of HAND has been previously suggested
124 [23-25]. Our earlier study of a small number of postmortem brain samples from HAND patients
125 found a trend towards increased abundance of APP and Tau relative to samples from uninfected
126 individuals [11]. To substantiate this finding, we now analyzed postmortem brain samples from
127 22 HIV-infected individuals diagnosed with HAND, 11 HIV-infected individuals without HAND

128 diagnosis, and 12 uninfected controls without diagnosed neurological disease (Table S1) for Tau
129 protein phosphorylated on Threonine 217 (p-Tau217). This Tau isoform has been shown to
130 strongly correlate with A β deposition [26] and was proposed as a marker for Alzheimer's disease
131 [20, 27, 28]. Results presented in Fig. 1A demonstrate that the abundance of p-Tau217, measured
132 by quantitative automated capillary Western blot, was significantly increased in samples from
133 HIV-infected individuals with HAND diagnosis, relative to HIV-infected individuals without
134 HAND diagnosis and especially relative to uninfected controls (supporting Western blot
135 evidence is presented in supplementary Fig. S1). Significant difference in the abundance of this
136 peptide was also observed between samples from HIV-infected individuals without HAND
137 diagnosis and uninfected controls (Fig. 1A). Given that p-Tau217 is an early marker of
138 neurodegeneration and Alzheimer's [20, 27, 28], these results are consistent with the role of
139 phosphorylated Tau in HAND pathogenesis and suggest that increase in this factor may precede
140 HAND diagnosis.

141 Amyloidogenic proteins and amyloid peptides are associated with lipid rafts [29-33], and
142 our previous study suggested that increased abundance of lipid rafts caused by Nef-containing
143 extracellular vesicles (exNef) may be the reason for the upregulation of APP and Tau [11].
144 Unfortunately, immunohistochemical analysis of lipid rafts in fresh-frozen tissue blocks was
145 technically challenging. We evaluated the abundance of lipid rafts in the brain tissue samples by
146 measuring the lipid raft marker flotillin 1 (by quantitative Western blot). Results of these
147 analyses are presented in Fig. S2. Although there was no difference in the abundance of flotillin
148 1 between brain samples from HIV-infected HAND-positive or HAND-negative individuals vs
149 uninfected individuals (Fig. S3), the abundance of p-Tau217 in HAND brain samples
150 significantly correlated with the abundance of flotillin 1 (Fig. 1B). This result is consistent with

151 the proposed relationship between lipid rafts and amyloid peptides [11, 32]. Of note, no such
152 correlation was observed in samples from uninfected brains, or brains from HIV-infected
153 individuals without HAND diagnosis (Fig. S4), suggesting that the relationship between rafts and
154 p-Tau217 formation is changed in HAND.

155 Results above suggest that abundance p-Tau217 in HAND brains may be associated with
156 the abundance of flotillin 1 and, by extension, of lipid rafts. Previous studies suggested that
157 downmodulation of ABCA1 in macrophages infected with HIV-1 or treated with exNef regulates
158 the abundance of lipid rafts [10-12, 34]. Our analysis showed a negative correlation between the
159 quantity of ABCA1 in all samples and the quantity of flotillin 1 (Fig. 1C). Comparison of
160 ABCA1 abundance between all samples from HIV-infected individuals to samples from
161 uninfected controls showed a trend towards reduced ABCA1 in the brains from infected
162 individuals, although the statistical significance was not achieved (Fig. 1D, supporting Western
163 blot evidence is in Fig. S5).

164 A known problem with detection of Nef in biological samples is that Nef proteins from
165 different HIV-1 strains have vastly varying ability to interact with different antibodies,
166 significantly affecting accuracy of comparison of the abundance of Nef between infected
167 participants. In addition, low specificity of most anti-Nef antibodies makes detection of low Nef
168 concentrations in clinical samples challenging. To get around these limitations, we detected Nef
169 in the brain samples by Western blot using four different antibodies raised against conservative
170 epitopes in different Nef regions: 3D12 mapped to the N-terminus region of HXB2 Nef, EH1
171 mapped to the C-terminus region, SN20 mapped to the SH3-binding domain of Nef, and 6JR
172 mapped to the C-terminus region (see Materials and Methods). Analysis with 3D12, SN20, and
173 EH1 antibodies was performed with all samples, except HIV 40, HIV 42, HIV 43, HAND 41,

174 HAND 44, HAND 45, and showed some cross-reactivity with cellular proteins in uninfected
175 samples (Fig. S6). All samples were analyzed using the 6JR antibody, which provided the
176 cleanest results, with least cross-reactivity with cellular proteins and negative results with all
177 uninfected samples (Fig. 2), and was used for the final interpretation of the Nef status. Table 1
178 summarizes results of this analysis. Eight of 22 HAND samples were Nef-positive, whereas only
179 one sample from HIV-positive individual without HAND diagnosis (HIV 19) was positive.

180 We further separated HAND samples into two groups, Nef-positive and Nef-negative,
181 and compared these groups for expression of ABCA1, flotillin 1, p-Tau217, and HAND severity
182 according to the clinical score (Fig. 3). Samples positive for Nef had lower abundance of
183 ABCA1 (Fig. 3A), higher abundance of flotillin (Fig. 3B), and higher abundance of p-Tau217
184 (Fig. 3C) relative to HAND samples with undetectable Nef. To determine whether the observed
185 differences were associated with the disease status, we compared the clinical HAND scores
186 between subjects providing the Nef-positive and Nef-negative brain samples. Clinical
187 neurological status was assessed using three existing dementia rating scales, American Academy
188 of Neurology (AAN), Memorial Sloan Kettering (MSK), and Frascati; the scales were not
189 universally applied to all individuals and in some cases disagreed (Table S1). Although a good
190 concordance between these scales has been reported, Frascati provided a more graded evaluation,
191 especially for mild cases [35]. We therefore used the Frascati score, where available, to grade the
192 clinical status, and the AAN score in the other cases (Table 2). The HAND clinical score was
193 significantly higher in individuals providing Nef-positive samples relative to individuals
194 providing samples with undetectable Nef (Fig. 3D), indicating that presence of Nef was
195 associated with a more severe disease. Multivariate regression analysis demonstrated that the
196 effect of detectable Nef on all continuous outcome variables (ABCA1, flotillin 1, p-Tau217)

197 considered jointly was significant in all HIV-infected individuals, in both the unadjusted model
198 and the adjusted model controlling for age, HIV duration, and viral load (Table S2). The
199 difference in results between analysis of all samples from HIV-positive individuals and those
200 from HAND only may be due to the smaller sample size in HAND only set.

201 Taken together, these results suggest that Nef, via modification of cholesterol metabolism
202 and lipid rafts may be one of the key causative factors promoting amyloid formation and disease
203 progression in HAND.

204

205 **Discussion**

206 The role of amyloid proteins in HAND pathogenesis remains an open question. In this
207 study, we demonstrated a significant increase in the brains of HIV-infected individuals of the
208 abundance of p-Tau217, which is also involved in pathogenesis of Alzheimer's disease [36].
209 This finding is consistent with our previous report where *in vitro* analysis suggested increased
210 abundance of amyloid precursor proteins and Tau in exNef-treated neuronal cells [11].
211 Surprisingly, this was true for brain samples from both HAND-diagnosed and HAND-free HIV-
212 infected individuals. One explanation is that HIV-infected individuals without HAND diagnosis
213 were at an early stage of disease when clinical manifestations could not yet be detected. This
214 explanation is consistent with a demonstrated utility of p-Tau217 as an early marker of
215 Alzheimer's, which allows detection of the disease at the stage preceding neurodegeneration and
216 clinical symptoms [37].

217 In these studies, we measured total flotillin 1 to evaluate the abundance of the lipid rafts.
218 Flotillins 1 and 2 are integral membrane proteins that assist in raft assembly [38-40]. While
219 flotillin 1 can be recruited to lipid rafts under certain stimulatory conditions [41], suggesting that

220 the number of flotillin molecules per lipid raft may vary, the level of flotillin expression, and
221 thus the per cell abundance of flotillin, appears to regulate the abundance of lipid rafts [42].
222 Therefore, the abundance of total flotillin 1 provides only a crude estimate of the abundance of
223 lipid rafts. This limitation is taken in the account in our interpretations.

224 Tau hyperphosphorylation is a characteristic feature of Alzheimer's disease, and a
225 number of studies established that phosphorylation of Tau in AD is closely linked to A β
226 pathology [26, 43, 44]. Lipid rafts were proposed to be the sites where Tau phosphorylation takes
227 place, and where phosphorylated Tau accumulates and interacts with A β during development of
228 Alzheimer's disease [33]. The observed correlation between abundancies of flotillin 1 and p-
229 Tau217 may reflect this paradigm, suggesting that lipid rafts may be a target for potential
230 therapeutic interventions in HAND, as had been proposed for other neurodegenerative diseases
231 [45]. We recognize, however, that, as discussed above, abundance of flotillin 1 provides only a
232 crude estimate of the abundance of lipid rafts, which may be the reason for low flotillin 1 in
233 some brain samples from subjects diagnosed with HAND, and high flotillin 1 in some samples
234 from HIV-infected individuals without the HAND diagnosis. But the most likely explanation for
235 this variability is that a driving factor in HAND pathogenesis is not the absolute amount, but the
236 change in the quality and quantity of the lipid rafts during disease progression. Our previous
237 studies documented functional impairment of lipid rafts associated with the effect of Nef [10],
238 which may occur without a major change in the raft abundance. We have previously reported
239 that exNef modify the fatty acid content of the lipid rafts [12], although the functional
240 consequences of these changes have not been investigated. Prospective studies will be needed to
241 evaluate this idea.

242 Our analysis demonstrated that majority of samples where Nef could be detected came
243 from individuals with diagnosed HAND. This result suggests an important role of Nef in HAND
244 pathogenesis. Indeed, Nef-positive samples were characterized by reduced ABCA1, increased
245 flotillin 1, increased p-Tau217, and increased clinical disease score. These results suggest that
246 Nef may drive ABCA1 downmodulation in HAND initiating lipid raft modifications and
247 subsequent pathological events.

248 A limitation of this study is a relatively small number of samples available for analysis.
249 Since samples were not cryopreserved, we could not perform flow cytometry analysis of the lipid
250 rafts, or cell-specific characterizations. Another limitation is lack of a sensitive quantitative assay
251 for Nef. Despite these limitations, our study supports the pathogenic role of Tau in HAND
252 pathogenesis, and suggests the sequence of events that lead to HAND: Nef EVs downmodulate
253 ABCA1 changing the properties of the lipid rafts, thus increasing formation of amyloid plaques
254 and Tau phosphorylation and fibrillation. Future studies in animal models will establish the
255 temporality of this ordering and will determine whether therapeutic treatments breaking the
256 pathogenic course described above can prevent development of HAND.

257

258 **Author Declarations**

259 *Ethics approval and consent to participate:* Not applicable. Anonymized samples were received
260 from repositories.

261 *Consent for publication:* All authors received the final version of the manuscript and agreed to
262 publish it.

263 *Availability of data and materials:* All the data are presented in the manuscript as the main or
264 supplementary material. The sources of materials are provided.

265 *Competing interests:* The authors declare no competing interests.

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268 *Authors' contributions:* TP performed the experiments and contributed to data analysis and
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270 interpretation, and manuscript preparation; MIB designed the experiments, contributed to results'
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277 samples from uninfected individuals. The brain samples from HIV-infected individuals with and
278 without HAND diagnosis were provided by The National NeuroAIDS Tissue Consortium
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281 Center (U24MH100930) National Neurological AIDS Bank (U24MH100929), California
282 NeuroAIDS Tissue Network (U24MH100928), Data Coordinating Center (U24MH100925).

283 **Compliance with Ethical Standards**

284 *Disclosure of potential conflicts of interest: The authors declare no conflicts of interest.*

285 *Research involving Human Participants and/or Animals:* Human samples analyzed in this study
286 were anonymized and were provided by NIH-maintained repositories.

287 *Informed consent:* Not applicable.

288

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Table 1. Results of Nef analysis in experimental samples.

Study ID	SN20	3D12	EH1	6JR
HAND 2	-	-	-	-
HAND 3	+	+	+	+
HAND 4	-	-	-	-
HAND 6	+	+	+	+
HAND 7	+	+	+	+
HAND 11	+	+	+	-
HAND 12	-	+	-	-
HAND 13	+	+	+	+
HAND 14	-	-	+	-
HAND 15	-	-	-	-
HAND 16	+	-	-	-
HAND 17	+	+	+	+
HAND 18	-	-	-	+
HAND 25	-	-	-	-
HAND 26	+	-	-	-
HAND 27	-	-	-	-
HAND 28	-	-	+	-
HAND 29	+	+	+	+
HAND 30	-	-	-	-
HAND 41	ND	ND	ND	+
HAND 44	ND	ND	ND	-
HAND 45	ND	ND	ND	-
HIV 1	-	-	-	-
HIV 5	-	-	-	-
HIV 8	+	-	-	-
HIV 9	-	-	-	-
HIV 10	+	+	-	-
HIV 19	+	+	+	+
HIV 20	+	-	-	-
HIV 24	-	-	-	-
HIV 40	ND	ND	ND	-
HIV 42	ND	ND	ND	-
HIV 43	ND	ND	ND	-
Uninf 21	+	-	-	-
Uninf 22	+	-	-	-
Uninf 23	+	-	-	-
Uninf 31	+	-	-	-
Uninf 32	-	-	-	-
Uninf 33	ND	ND	ND	-
Uninf 34	-	-	-	-
Uninf 35	+	-	-	-
Uninf 36	-	-	-	-
Uninf 37	+	-	-	-
Uninf 38	-	-	ND	-
Uninf 39	+	-	+	-

ND – not done. + - Nef detected, - - Nef undetected.

Table 2. HAND severity evaluation in donors of post-mortem brain samples^A

ID	Frascati score	AAN score	HAND score in the study
HAND 2		3	3
HAND 3		5	5
HAND 4		4	4
HAND 6		4	4
HAND 7		5	5
HAND 11		3	3
HAND 12		2	2
HAND 13		5	5
HAND 14		3	3
HAND 15		3	3
HAND 16	3		3
HAND 17	3		3
HAND 18	2		2
HAND 25	3		3
HAND 26		3	3
HAND 27		3	3
HAND 28		3	3
HAND 29		3	3
HAND 30		3	3
HAND 41	5	4	5
HAND 44	5	4	5
HAND 45	3	5	3

^AHAND score used in the study was based on Frascati score where available, and AAN score in other cases.

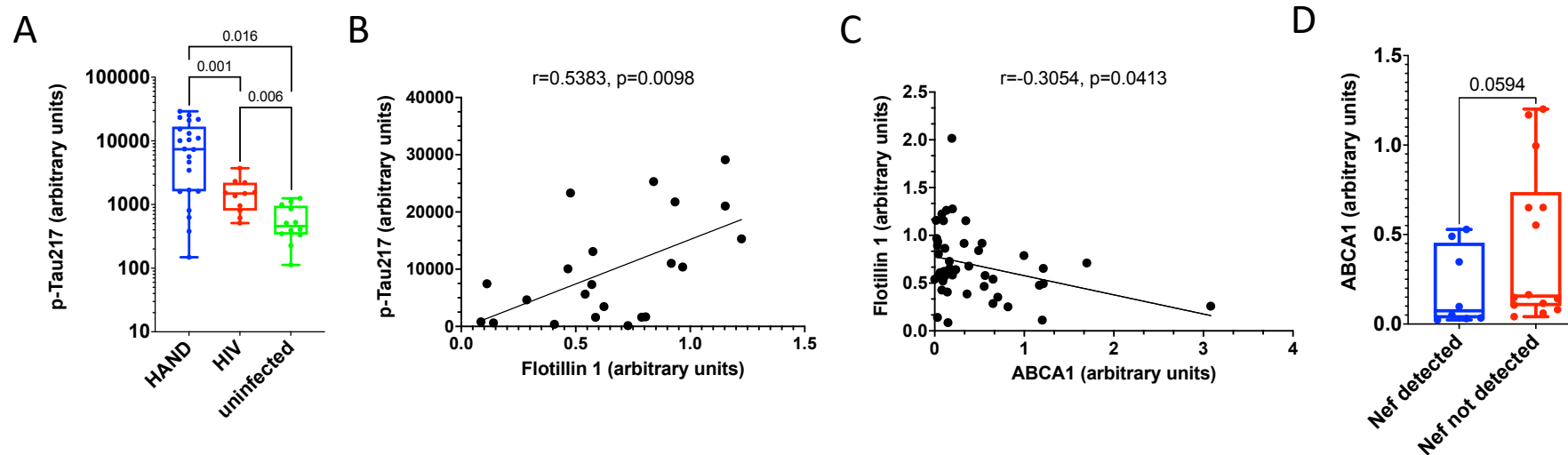


Figure 1. Analysis of pathogenic correlates of HAND. A – Analysis of p-Tau217 in brain samples from uninfected individuals and HIV-infected individuals with and without HAND. Data points representing p-Tau217 adjusted to total protein were obtained using ProteinSimple Compass software and are presented as arbitrary units. Group comparisons were made by Kruskal-Wallis test with post hoc Dwass, Steel, Critchlow-Fligner multiple comparison analysis adjustment for FWER (DSCF option in SAS NPAR1WAY procedure). B – Simple linear regression analysis of p-Tau217 and Flotillin 1 in HAND samples (data adjusted to total protein obtained using Compass software). C - Simple linear regression analysis of ABCA1 and Flotillin 1 in all samples (protein abundance adjusted to total protein was obtained using Compass software). D – Analysis of ABCA1 in HIV-infected (HAND-positive and HAND-negative samples) vs uninfected samples. Data points representing ABCA1 abundance adjusted to total protein were obtained using Compass and are presented as box and whiskers plot with p value calculated by Mann Whitney nonparametric two-tailed t test.

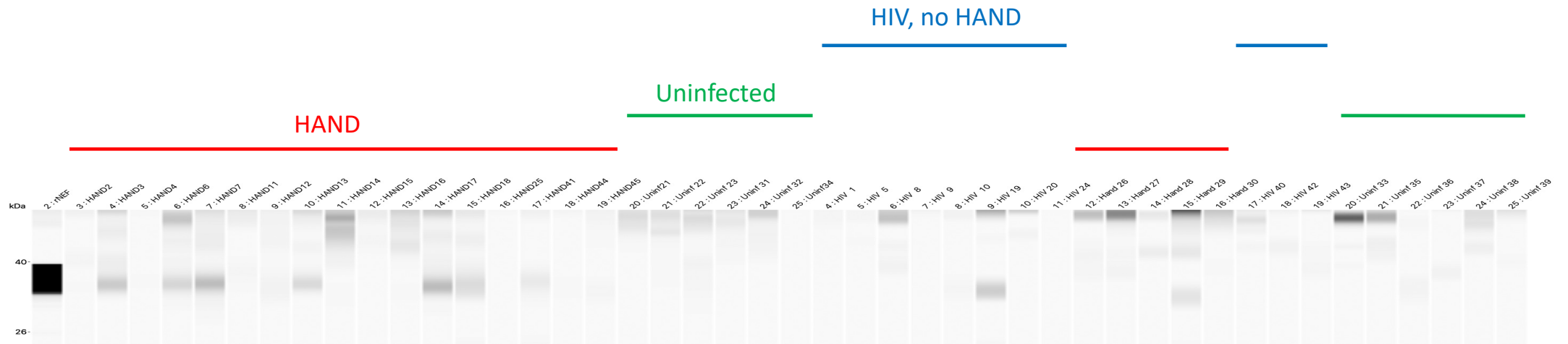


Figure 2. Analysis of Nef in brain samples. Brain lysates were assayed by Western immunoblot using the 6JR mouse monoclonal anti-Nef antibody from Abcam. Samples were run on ProteinSimple Jess instrument and analyzed by Compass software. Color-coded lines denote group assignment of the samples.

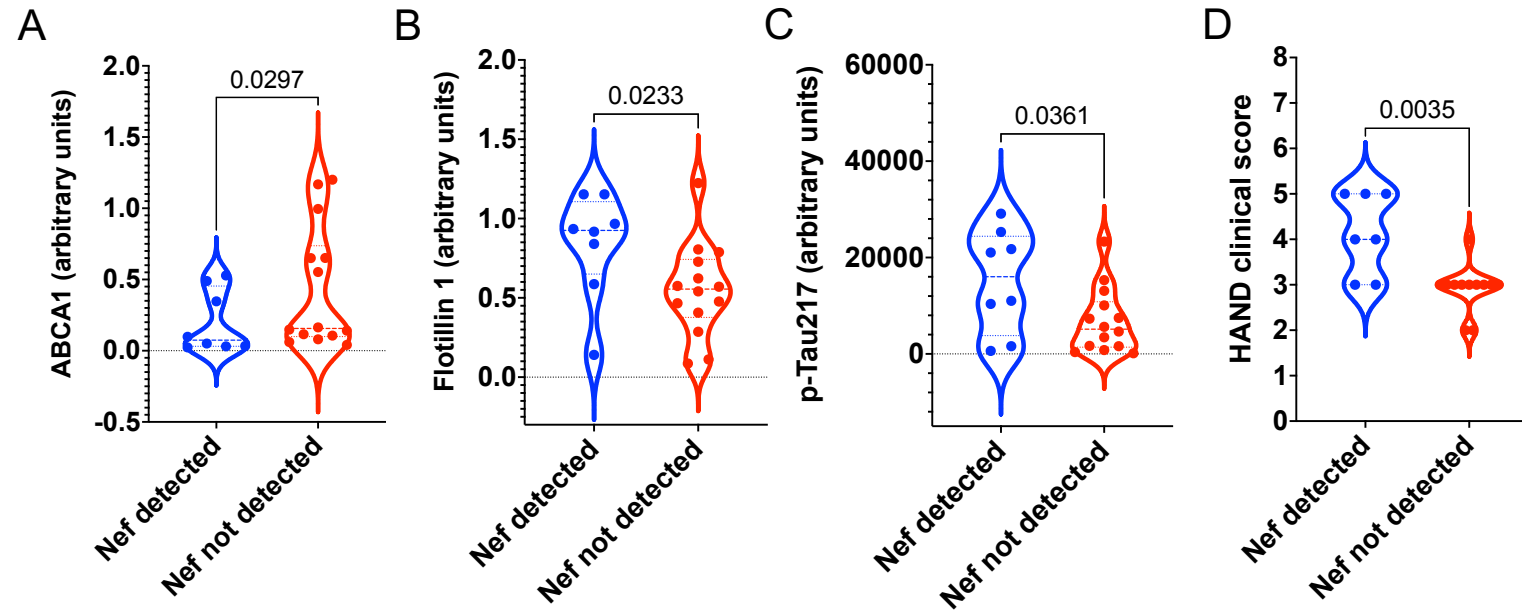


Figure 3. Comparative analysis of Nef-positive and Nef-negative brain samples from HAND-diagnosed individuals. A, B, C - Data points for ABCA1 (A), Flotillin 1 (B), and p-Tau217 (C) adjusted to total protein levels were obtained using ProteinSimple Compass software and are presented as arbitrary units. Results are presented as violin plots, showing p values calculated for unpaired parametric two-tailed t test. D – HAND clinical scores are presented for individuals from whom Nef-positive and Nef-negative brain samples were obtained as violin plots, showing p values calculated for unpaired non-parametric Mann-Whitney two-tailed t test.