Supplementary Data

Aftins Increase Amyloid- β_{42} , Lower Amyloid- β_{38} , and Do Not Alter Amyloid- β_{40} Extracellular Production *in vitro*: Toward a Chemical Model of Alzheimer's Disease?

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CHEMISTRY

Preparation of Aftins-1 (3) and -2 (4)

The synthesis of Aftins MRT2-309 (2), Aftin-1 (3) and Aftin-2 (4) is depicted in Scheme 1.

N6-Benzyl-N6-methyl-9H-purine-2,6-diamine (2)

N-Methylbenzylamine and NEt₃ were added to a solution of 2-amino-6-chloropurine in *n*-butanol. The mixture was stirred at 85°C for 6 h. After cooling to 20°C, Aftin-2 separated upon filtration of the precipitate. Yield 74%, mp 130°C. ¹H-NMR (CDCl₃) δ ppm: 3.35 (brs 3H); 4.75 and 5.25 (2 brs, CH₂); 7.35 (m, 5H, C₆H₅); 7.65 (s, 1H, 8-H).

N6-Benzyl–9-isopropyl-N6-methyl-purine-2,6 -diamine (Aftin-1) (3)

Yield 72%, mp 110°C. ¹H-NMR (CDCl₃) δ ppm: 1.45 (d, 6H, CH (CH₃)₂); 3.30 (brs, CH₃N); 4.51 (brs CH₂); 4.62 (hept, 1H, CH (CH₃)₂); 5.23 (brs, NH₂).7.2 (m, 5H, C₆H₅) 7.48 (s, 1H, 8-H). ¹³C-NMR 22.74 (CHCH₃); 35.83 (brs, CH₃N); 45.76 (CH); 53.26 (CH₂N); 115.25 (CH); 127.07 (CH); 127.65 (CH); 128.49 (CH) 129.05; 133.53 (CH); 138.33 (CH); 152.54 (CH); 155.27 (CH); 159.08 (CH).

N6-Benzyl–9-cyclopentyl-N6-methyl-purine-2,6 -diamine (Aftin-2) (4)

Mp 160°C. ¹H NMR (CDCl₃) 1.70 (m, 2H, cyclopentyl); 1.83 (m, 4H, cyclopentyl); 2.16 (m, 2H, cyclopentyl); 3.28 (brs, 3H, CH₃); 4.61 (brs, 2H, CH₂-C₆H₅); 4.73 (quint, 1H, cyclopentyl); 4.61 (brs, 2H, CH₂); 5.21 (brs, NH₂); 7.25 (m, 5H); 7.46 (s, 1H).

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Scheme 1. Reagents and conditions. a) N-methylbenzylamine, N (Et)₃, in n-butanol, 85°C; b) *iso*-propylbromide, K₂CO₃, DMSO 16–18°C; c) 2-bromocyclopentane, K₂CO₃, DMSO 16–18°C.



Scheme 2. Reagents and conditions. a) NaH, benzylic alcohol; b) (R)-2-aminobutanol, 140°C.

Preparation of Aftin-3 (5)

The synthesis of Aftin-3 (5) is presented in Scheme 2. Compound 5a was prepared as described [1].

6-Benzyloxy-2-chloro-9-isopropyl-purine (5b)

THF (20 mL) was slowly added under nitrogen to NaH 60% (1.70 g; 54.6 mmol) dispersion in mineral oil cooled by an external bath. Benzyl alcohol (2 mL, 18.5 mmol) in THF (10 mL) was added dropwise at 0°C. After 15 min stirring at the same temperature, 2, 6-dichloro-9-*iso*-propylpurine, 5a (3.64 g, 15.8 mmol) in 20 mL THF was slowly added. After 12 h stirring 20°C, the mixture was cooled at 0°C and 100 mL 2M NH₄Cl solution was slowly added. The mixture was extracted by AcOEt (3×20 mL) and the organic layer was washed with water (2×20 mL). After drying (Na₂SO₄) 5b crystallized upon concentration of the solution. ¹H-NMR (CDCl₃) δ ppm: 1.63 (d, 6H, J=7.2 Hz), 4.85 (hept, 1H, J=7.2 Hz), 5.60 (s, 2H), 7.32–7.41 (m, 3H), 7.52 (d, 2H, J=6.8 Hz), 8.82 (s, 1H, 8-H).

(2R)-2-[(6-Benzyloxy-9-isopropyl-purin-2-yl) amino]butan-1-ol (Aftin-3) (5)

A mixture of 5b (1.45 g, 4.7 mmol) and (R)-2-aminobutanol (3.58 mL, 0.4 mmol) was heated 6 h at 140°C. After cooling, the mixture was extracted with CH₂Cl₂ washed with

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Scheme 3. Reagents and conditions. a) N-methylbenzylamine, N (Et)₃, in n-butanol, 85°C; b) *iso*-propylbromide, K₂CO₃, DMSO 16–18°C; c) (R)-2-aminobutanol, 160°C.



Scheme 4. Reagents and conditions. a) N-methylaniline, N (Et)₃, in n-butanol, $95^{\circ}C$; b) isopropylbromide, K₂CO₃, DMSO 16–18°C; c) (R)-2-aminobutanol, $160^{\circ}C$.

water (2×20 mL). Derivative 5 crystallized upon trituration with AcOEt. ¹H- NMR (CDCl₃) δ ppm: 0.99 (t, 3 H, *J* = 7.2 Hz), 1.53–1.61 (m, 1 H), 1.63 (d, 6 H, *J* = 7.2 Hz), 1.74–1.80 (m, 1 H), 3.59–3.69 (m, 2 H), 4.00–4.08 (m, 1 H), 4.85 (septet, 1H, *J* = 7.2 Hz), 5.60 (s, 2 H), 7.32–7.41 (m, 3 H), 7.52 (d, 2 H, *J* = 6.8 Hz), 8.82 (s, 1 H).

Preparation of Aftin-4 (6)

The preparation of Aftin-4 (6) is depicted in Scheme 3.

The amination of position 6 was first performed upon heating 2,6-dichloropurine with Nmethylbenzylamine in *n*-BuOH. Alkylation of 6a was achieved using an alkylhalide in a dipolar aprotic solvent such as DMSO. In the last step, the amination was achieved upon heating the chloro derivatives 6b with an amine (Scheme 3).

N-Benzyl-2-chloro-N-methyl-9H-purin-6amine (6*a*)

N-Methylbenzylamine (10.5 mL, 79.36 mmol) and triethylamine (29 mL, 211.63 mmol) were added to a solution of 2,6-dichloropurine (10.0 g, 52.91 mmol) in n-butanol (80 mL). The mixture was heated at 100° C for 3 h. The reaction was cooled to 40° C, and the

precipitate was filtered at this temperature, washed with 10 mL cold (15° C) H₂O, and dried overnight in vacuo.

N-Benzyl-2-chloro-9-isopropyl-N-methyl-purin-6-amine (6b)

2-Bromopropane (25.77 mL, 275.37 mmol) was added to a solution of 1 (10.76 g, 39.34 mol) and K₂CO₃ (21.74 g, 157.35 mmol) in 80 mL DMSO. After 12 h stirring, the mixture was diluted with 200 mL H₂O and extracted with AcOEt (3×20 mL). After concentration the remaining solid was triturated with 5 mL AcOEt filtered and dried overnight in vacuo. Yield 65%, mp 103–106°C. ¹H-NMR (DMSOd6) δ ppm: 1.47 (d, 6H, (CH₃)₂CH); 3.12 and 3.60 (2 brs, 3H, CH₃N); 4.75 (hept, 1H, *J*=Hz) 4.90 and 5.50 (2 brs, 2H, CH₂C₆H₅); 75 (s, 1H, 8-H). ¹³C-NMR δ ppm: 22.7, 46.62, 119.07, 127.46, 128.64; 136.12; 137.31; 153.65.

(2R)-2-[Amino-[6-(benzyl(methyl)amino)-9isopropyl-purin-2-yl]amino]butan-1-ol (Aftin-4) (6)

A mixture of the chloro derivative 2 (12.0 g, 37.99 mmol) (R)-2-aminobutanol (28.47 mL, 304 mmol) was heated 4 h at 160°C. After cooling, the mixture was extracted with CH_2Cl_2 washed with water



Scheme 5. Reagents and conditions. a) BrEt, K_2CO_3 , DMF, $20^{\circ}C$; b) Fe, HCl, NH₄Cl, EtOH; c) 1. C_6H_5CONCO , 2. NaOH $80^{\circ}C$; d) K_2CO_3 , DMSO, $85^{\circ}C$; e) Br₂, HBr, CH₃COOH; f) DMF 75^{\circ}C.

(2×20 mL). Derivative 3 was crystallized upon trituration with AcOEt, followed by recrystallisation from 2-propanol. Yield, 65%, mp 114–117°C. ¹H-NMR (DMSO_{d6}) δ ppm:1.01 (t, 3H, CH₃CH₂); 1.49 (d, 6 H, (CH₃)₂CH); 1.51 (m, 2 H, CH₂CH₃); 3.26 (brs, 3 H, CH₃N) 3.77 and 3.81 (2 m, 2 H, CH₂O); 3.78 (m 1H, CHO); 4.55 (m, hept, 1 H), 4.70 (d, 1 H, NH); 5;21 (brs, CH₂C₆H₅); 7.28 (m, 5 H, C₆H₅); 7.45 (s, 1 H, 8-H). ¹³C-NMR δ ppm: 10.97; 22.52; 22, 56; 25.08; 46.16; 53.38 56.33; 68.76; 115.11; 127.12; 127.58; 128.52; 133.36; 138.19; 151.84, 155.08; 159.44.

Preparation of Aftin-5 (7)

The preparation of Aftin-5 (7) is depicted in Scheme 4. Briefly the amination of 2, 6-dichloropurine was first performed upon heating with N-methylaniline and NEt₃ in *n*BuOH. Alkylation was then performed followed by amination at the 2 position. This last step was achieved upon heating until completion with an 8 fold excess of (R)-2-aminobutanol at 160° C.

(2R)-2-[[9-isopropyl-6-(N-methylanilino)purin-2-yl]amino]butan-1-ol (Aftin-5) (7)

Yield 63%. ¹H-NMR (CDCl₃) δ ppm: 0.92 (t, 3 H, CH₃); 1.45 (d, 6 H, (CH₃)₂CH); 1.49 (m, 2 H,

CH₂CH₃); 3.50 (m, 1 H, CHN); 3.69 (m, 2 H CH₂-O); 3.72 (s, 3 H, CH₃N); 4.54 (hept, 1 H, (CH₃)₂CH); 4.79 (brs, 2 H, NH₂); 7.24 and 7.34 (2 m, 5 H, C₆H₄) 7.39 (s, 1 H, 8-H). ¹³C-NMR: 10.94 (CH₃); 22.48 and 22.54 (2 CH₃); 24.97 (CH₂); 40.21 (CH); 46.13 (CH); 56.28 (CH); 68.37 (CH₂); 115.3 (C); 126.43 (CH); 126.99 (CH); 129.04 (CH); 134.33 (CH) 14.28 (C); 154.13 (C); 155.14 (C); 159.04 (C).

Preparation of the 'Torrey Pines' compound (9)

The synthesis of the 'Torrey Pines' compound 9 was achieved as depicted in Scheme 5.

N1, N1-diethyl-4-methyl-N3-[4-[4-(4-methylimidazol-1-yl)phenyl]thiazol-2yl]benzene-1, 3-diamine (9)

¹H-NMR (CDCl₃) δ ppm: 1.05 (t, 6 H, 2CH₃CH₂); 2.10 (s, 3 H, CH₃); 2.18 (s, 3 H, CH₃); 3.40 (m, 4 H, 2CH₂); 6.45 (dd, J = 4 Hz, 2 Hz, 1 H); 6.98 (m, 2 H); 7.35 (d, 2 H, J = 7 Hz); 7.75 (s, 1 H, imidazol); 7.92 (d, 2 H, J = 7 Hz). ¹³C-NMR (DEPT135): 12.67 (CH₃); 13.74 (CH₃); 16.57 (CH₃); 44.60 (CH₂); 102.13 (CH); 104.00 (CH); 108.32 (CH); 114.48 (CH); 121.00 (CH); 127.37 (CH); 131.67 (CH); 134.51 (CH). KINASE

CAMK1D

CAMK1G

CAMK2A

CAMK2B

CAMK2D

CAMK2G

SELECTIVITY STUDIES

Supplementary Table 1

Discoverx KinomeScan Selectivity Panel (442 kinases). Enzymes were prepared and interactions assays were run in the presence of 10 μ M Aftin-5, as described in [2]. A semi-quantitative scoring of this primary screen was estimated. This score relates to a probability of a hit rather than strict affinity. Scores >10, between 1–10 and <1 indicate the probability of a being a false positive is <20%,<10%, <5 %, respectively

KINASE	SCORE	CAMK4
AK1	98	CAMKKI
ABL1 (E255K)-phosphorylated	89	CASK
ABL1 (F317I)-nonphosphorylated	72	CASK CDC2L1
ABL1 (F317I)-phosphorylated	100	CDC2L1 CDC2L1
ABL1 (F317L)-nonphosphorylated	100	CDC2L2 CDC2L5
ABL1 (F317L)-phosphorylated	74	CDC2L5
ABL1 (H396P)-nonphosphorylated	100	CDKII
ABL1 (H396P)-phosphorylated	70	CDK2
ABL1 (M351T)-phosphorylated	50	CDK3
ABL1 (0252H)-nonphosphorylated	100	CDK4-cyclinD1
ABL 1 (Q252H)-nhosphorylated	96	CDK4-cyclinD3
ABL1 (T315I)-nonphosphorylated	100	CDK5
ABL 1 (T3151) phosphorylated	52	CDK7
ABL1 (15151)-phosphorylated	52	CDK8
ABL1 (1233F)-phosphorylated	00 87	CDK9
ADD 1 nhosphorylated	0/	CDKL1
ADL1-phosphorylated	91	CDKL2
	100	CDKL3
	100	CDKL5
CVRIB	100	CHEK1
ACVR2A	100	CHEK2
ACVR2B	100	CIT
ACVRL1	100	CLK1
ADCK3	100	CLK2
ADCK4	63	CLK3
AKT1	92	CLK4
AKT2	100	CSF1R
AKT3	91	CSK
ALK	94	CSNK1A1
MPK-alpha1	100	CSNK1A1L
MPK-alpha2	91	CSNK1D
NKK1	87	CSNK1E
ARK5	100	CSNK1G1
ASK1	83	CSNK1G2
ASK2	93	CSNK1G3
AURKA	84	CSNK2A1
URKB	89	CSNK2A2
AURKC	100	CTK
AXL	86	DAPK1
BIKE	100	DAPK2
BLK	80	DAPK3
3MPR1A	72	DCAMKI 1
3MPR1B	70	DCAMEL 2
3MPR2	86	DCANIKL2
BMX	93	DUCAWIKLS
BRAF	100	DDR1
3RAF (V600E)	100	DDK2
SRK	98	DLK
BRSK1	66	DMPK
RRSK2	86	DMPK2
110124	00	DRAK1
8TK	100	DD IIIA
TK AMK 1	100	DRAK2

Supplementary Table 1	
(Continued)	

SCORE

Supplementary Table 1 (Continued)		Supplementary Table 1 (Continued)		
KINASE	SCORE	KINASE	SCORE	
DYRK2	79	GRK4	98	
EGFR	100	GRK7	89	
EGFR (E746-A750del)	95	GSK3A	91	
EGFR (G719C)	94	GSK3B	72	
EGFR (G719S)	94	HCK	92	
EGFR (L747-E749del, A750P)	86	HIPK1	99	
EGFR (L747-S752del, P753S)	79	HIPK2	91	
EGFR (L747-T751del, Sins)	94	HIPK3	39	
EGFR (L858R)	93	HIPK4	78	
EGFR (L858R, 1790M)	100	HPK1	84	
EGFR (L861Q)	75	HUNK	82	
EGFR (5/52-1/59del)	/9		96	
EUFK (1/90M) $EIE2AV1$	100	IOFIR IKK alpha	100	
	90	IKK-aipila IKK beta	100	
	90	IKK-ocia IKK-ensilon	100	
EPHA3	86	INSR	75	
EPHA4	91	INSRR	95	
EPHA5	100	IRAK1	59	
EPHA6	100	IRAK3	60	
EPHA7	100	IRAK4	54	
EPHA8	90	ITK	100	
EPHB1	100	JAK1 (JH1domain-catalytic)	82	
EPHB2	100	JAK1 (JH2domain-pseudokinase)	99	
EPHB3	100	JAK2 (JH1domain-catalytic)	95	
EPHB4	100	JAK3 (JH1domain-catalytic)	100	
EPHB6	65	JNK1	92	
ERBB2	92	JNK2	100	
ERBB3	100	JNK3	100	
ERBB4	96	KIT	88	
ERK1	100	KIT (A829P)	100	
ERK2	100	KIT (D816H)	75	
ERK3	94	KIT (D816V)	91	
EKK4	12	KII (L5/6P) KIT (1/550D)	/9	
	98	KII (V559D) KIT (V559D, T670I)	80 100	
ERN0 FRN1	100 97	KIT (V559D, V654A)	100	
FAK	91	LATS1	100	
FER	100	LATS2	84	
FES	100	LCK	100	
FGFR1	100	LIMK1	100	
FGFR2	98	LIMK2	100	
FGFR3	81	LKB1	100	
FGFR3 (G697C)	84	LOK	91	
FGFR4	100	LRRK2	100	
FGR	100	LRRK2 (G2019S)	100	
FLT1	100	LTK	100	
FLI3	59	LYN	100	
FLT3 (D835H)	99	LZK	100	
FLI3 (D833 I)	98	MAK MAD2K1	95	
FLI3(IID) $FLT3(K662O)$	100		90	
FLT3 (N8411)	84	MAP3K15 MAP3K2	00	
FLT3 (R834O)	84	MAP3K3	100	
FLT4	100	MAP3K4	100	
FRK	100	MAP4K2	90	
FYN	100	MAP4K3	100	
GAK	100	MAP4K4	100	
GCN2 (Kin.Dom.2, S808G)	100	MAP4K5	100	
GRK1	81	MAPKAPK2	100	

Supplementary Table 1

Supplementary Table 1	
(Continued)	

Supplementary Table 1 (Continued)

KINASE	SCORE	KINASE	SCORE
MAPKAPK5	100	PAK4	90
MARK1	98	PAK6	100
MARK2	94	PAK7	86
MARK3	92	PCTK1	82
MARK4	84	PCTK2	82
MAST1	100	РСТКЗ	100
MEK1	100	PDGFRA	100
MEK2	100	PDGFRB	100
MEK3	97	PDPK1	95
MEK4	100	PFCDPK1 (P. falciparum)	100
MEK5	91	PFPK5 (P falcinarum)	84
MEK6	100	PFTAIRE2	100
MELK	100	PFTK1	100
MERTK	100	PHKG1	96
MET	89	PHKG2	100
MET (M1250T)	100	PIK3C2B	100
MET (Y1235D)	88	PIK3C2G	100
MINK	100	PIK3CA	100
MKK7	100	PIK3CA (C420R)	100
MKNK1	89	PIK3CA (E542K)	97
MKNK2	92	PIK3CA (E545A)	100
MICK	100	PIK3CA (E545K)	100
MLCK MLK1	03	PIK3CA (H1047I)	100
MLK1 MLK2	90	PIK3CA (H1047E)	100
MLK2	93	PIK3CA (18001)	100
MRCKA	98	PIK3CA (M10/3I)	100
MACKA	100	$\mathbf{D}\mathbf{K}3\mathbf{C}\mathbf{A} \left(\mathbf{O}546\mathbf{K}\right)$	86
MIRCIND MST1	100	DIK3CB	100
MSTID	100	PIK3CD	100
MST1IK MST2	90 60	PIK3CG	100
MST2	100	DIVACE	62
MST4	100	FIK4CB DIM1	03
MIS14 MTOP	100	PIMI DIM2	94
MUSK	100	DIM2	94
MUSK MVI V	72		98
MILK MVLV2	100	DID5V1C	99
MYI KA	100	DID5V2B	100
MYO3A	100	DID5K2C	100
MYO2P	100	$DK \Delta C$ alpha	100
	100	PKAC bata	04 100
	100	FKAC-UCIA	100
NDR2 NEV1	97	F KWI I I I DVN1	100
	00	F KINI DVN2	100
NEK1	98	PKINZ DKNB (M. tubaraulasis)	89 100
NEK2	97	PKIND (M. IUDERCUIOSIS) DI K 1	100
NEKS NEV4	100		100
NEK5	100		07 100
NEKJ	80		100
NEKO	100	PLK4	63 100
NEK/	94	PKKCD	100
	100	PKKCE	100
	88	PKKCH	100
	90		100
	45		100
pso-appna	100		81
	100		87
	100		84
p38-gamma	100	PKKG1	92
PAKI	73	PKKG2	99
PAK2	100	PKKK	90
PAK3	100	PKKX	100

KINASE SCORE KINASE PRP4 88 STK36 PYK2 100 STK39 QSK 98 SYK RAF1 100 TAK1 RET 99 TAOK1 RET (M918T) 89 TAOK2 RET (W804L) 95 TAOK3 RET (V804L) 95 TAOK3 RET (V804M) 94 TBK1 RIOK1 96 TEC RIOK2 100 TESK1 RIOK3 87 TGFBR1 RIPK1 89 TGFBR2 RIPK2 84 TIE1 RIPK5 77 TLK1 ROCK1 92 TLK2 ROCK1 92 TLK2 ROCK2 100 TNK1 RPS6KA4 (Kin.Dom.1-N-terminal) 93 TNK2 RPS6KA5 (Kin.Dom.2-C-terminal) 100 TRKA RPS6KA5 (Kin.Dom.2-C-terminal) 100 TRKA RPS6KA5 (Kin.Dom.2-C-terminal) 100	CODE
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PRP4 88 STK36 PYK2 100 STK39 QSK 98 SYK RAF1 100 TAK1 RET 99 TAOK1 RET (M918T) 89 TAOK2 RET (V804L) 95 TAOK3 RET (V804M) 94 TBK1 RIOK1 96 TEC RIOK2 100 TESK1 RIOK3 87 TGFBR1 RIPK4 100 TE2 RIPK4 100 TE2 RIPK4 100 TIE2 RIPK5 77 TLK1 ROCK1 92 TLK2 ROS1 100 TNIK ROS1 100 TNIK RPS6KA4 (Kin.Dom.1-N-terminal) 93 TNK2 RPS6KA5 (Kin.Dom.1-N-terminal) 100 TRKA RPS6KA5 (Kin.Dom.2-C-terminal) 100 TRKA RPS6KA5 (Kin.Dom.2-C-terminal) 86 TRKC	SCORE
PYK2 100 STK39 QSK 98 SYK RAF1 100 TAK1 RET 99 TAOK1 RET (M918T) 89 TAOK2 RET (W94L) 95 TAOK3 RET (V804L) 95 TAOK3 RET (V804M) 94 TBK1 RIOK1 96 TEC RIOK2 100 TESK1 RIOK3 87 TGFBR1 RIPK1 89 TGFBR2 RIPK2 84 TIE1 RIPK4 100 TIE2 RIPK5 77 TLK1 ROCK1 92 TLK2 ROCK1 92 TLK2 ROS1 100 TNK RPS6KA4 (Kin.Dom.1-N-terminal) 93 TNK2 RPS6KA5 (Kin.Dom.1-N-terminal) 80 TNN13K RPS6KA5 (Kin.Dom.1-N-terminal) 100 TRKA RPS6KA5 (Kin.Dom.1-N-terminal) 100 TRKA RPS6KA5 (Kin.Dom.1-N-terminal) 86 TRKC	100
QSK 98 SYK RAF1 100 TAK1 RET 99 TAOK1 RET (M918T) 89 TAOK2 RET (V804L) 95 TAOK3 RET (V804M) 94 TBK1 RIOK1 96 TEC RIOK2 100 TESK1 RIOK3 87 TGFBR1 RIPK1 89 TGFBR2 RIPK2 84 TIE1 RIPK3 77 TLK1 ROCK1 92 TLK2 ROCK2 100 TNIK ROS1 100 TNK1 RPS6KA4 (Kin.Dom.1-N-terminal) 93 TNK2 RPS6KA5 (Kin.Dom.1-N-terminal) 80 TNN13K RPS6KA5 (Kin.Dom.2-C-terminal) 100 TRKA RPS6KA5 (Kin.Dom.2-C-terminal) 100 TRKB RSK1 (Kin.Dom.2-C-terminal) 86 TRKD	100
RAF1 100 TAK1 RET 99 TAOK1 RET (M918T) 89 TAOK2 RET (V804L) 95 TAOK3 RET (V804M) 94 TBK1 RIOK1 96 TEC RIOK2 100 TESK1 RIOK3 87 TGFBR1 RIPK1 89 TGFBR2 RIPK2 84 TIE1 RIPK4 100 TIE2 RIPK5 77 TLK1 ROCK1 92 TLK2 ROCK2 100 TNIK ROS1 100 TNK1 RPS6KA4 (Kin.Dom.1-N-terminal) 93 TNK13K RPS6KA5 (Kin.Dom.1-N-terminal) 100 TRKA RPS6KA5 (Kin.Dom.1-N-terminal) 86 TRKC	100
RET 99 TAOK1 RET (M918T) 89 TAOK2 RET (V804L) 95 TAOK3 RET (V804M) 94 TBK1 RIOK1 96 TEC RIOK2 100 TESK1 RIOK3 87 TGFBR1 RIPK1 89 TGFBR2 RIPK2 84 TIE1 RIPK4 100 TIE2 RIPK5 77 TLK1 ROCK1 92 TLK2 ROCK1 92 TLK2 ROCK1 93 TNK2 ROS1 100 TNIK RPS6KA4 (Kin.Dom.1-N-terminal) 93 TNK2 RPS6KA5 (Kin.Dom.1-N-terminal) 100 TRKA	91
RET (M918T) 89 TAOK2 RET (V804L) 95 TAOK3 RET (V804M) 94 TBK1 RIOK1 96 TEC RIOK2 100 TESK1 RIOK3 87 TGFBR1 RIPK1 89 TGFBR2 RIPK2 84 TIE1 RIPK4 100 TIE2 RIPK5 77 TLK1 ROCK1 92 TLK2 ROS1 100 TNIK RPS6KA4 (Kin.Dom.1-N-terminal) 93 TNK12 RPS6KA5 (Kin.Dom.1-N-terminal) 100 TRKA RPS6KA5 (Kin.Dom.1-N-terminal) 100 TRKA RPS6KA5 (Kin.Dom.1-N-terminal) 86 TRKC	97
RET (V804L) 95 TAOK3 RET (V804M) 94 TBK1 RIOK1 96 TEC RIOK2 100 TESK1 RIOK3 87 TGFBR1 RIPK1 89 TGFBR2 RIPK2 84 TIE1 RIPK4 100 TIE2 RIPK5 77 TLK1 ROCK1 92 TLK2 ROS1 100 TNIK RPS6KA4 (Kin.Dom.1-N-terminal) 93 TNNI3K RPS6KA5 (Kin.Dom.2-C-terminal) 100 TRKA RPS6KA5 (Kin.Dom.2-C-terminal) 86 TRKC	100
RET (V804M) 94 TBK1 RIOK1 96 TEC RIOK2 100 TESK1 RIOK3 87 TGFBR1 RIPK1 89 TGFBR2 RIPK2 84 TIE1 RIPK4 100 TIE2 RIPK5 77 TLK1 ROCK1 92 TLK2 ROS1 100 TNIK RPS6KA4 (Kin.Dom.1-N-terminal) 93 TNNI3K RPS6KA5 (Kin.Dom.2-C-terminal) 100 TRKA RPS6KA5 (Kin.Dom.2-C-terminal) 86 TRKD	88
RIOK1 96 TEC RIOK2 100 TESK1 RIOK3 87 TGFBR1 RIPK1 89 TGFBR2 RIPK2 84 TIE1 RIPK4 100 TIE2 RIPK5 77 TLK1 ROCK1 92 TLK2 ROCK1 92 TLK2 ROS1 100 TNK1 RPS6KA4 (Kin.Dom.1-N-terminal) 93 TNK2 RPS6KA5 (Kin.Dom.1-N-terminal) 100 TRKA RPS6KA5 (Kin.Dom.2-C-terminal) 100 TRKA RPS6KA5 (Kin.Dom.2-C-terminal) 86 TRKD	74
RIOK2 100 TESK1 RIOK3 87 TGFBR1 RIPK1 89 TGFBR2 RIPK2 84 TIE1 RIPK4 100 TIE2 RIPK5 77 TLK1 ROCK1 92 TLK2 ROCK1 92 TLK2 ROS1 100 TNIK RPS6KA4 (Kin.Dom.1-N-terminal) 93 TNK1 RPS6KA5 (Kin.Dom.1-N-terminal) 100 TRKA RPS6KA5 (Kin.Dom.2-C-terminal) 100 TRKA RPS6KA5 (Kin.Dom.2-C-terminal) 86 TRKD	96
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RIPK1 89 TGFBR2 RIPK2 84 TIE1 RIPK4 100 TIE2 RIPK5 77 TLK1 ROCK1 92 TLK2 ROCK2 100 TNIK ROS1 100 TNK1 RPS6KA4 (Kin.Dom.1-N-terminal) 93 TNK2 RPS6KA5 (Kin.Dom.2-C-terminal) 80 TNNI3K RPS6KA5 (Kin.Dom.2-C-terminal) 100 TRKA RPS6KA5 (Kin.Dom.2-C-terminal) 86 TRKD	99
RIPK2 84 TIE1 RIPK4 100 TIE2 RIPK5 77 TLK1 ROCK1 92 TLK2 ROCK2 100 TNIK ROS1 100 TNK1 RPS6KA4 (Kin.Dom.1-N-terminal) 93 TNK2 RPS6KA4 (Kin.Dom.2-C-terminal) 80 TNN13K RPS6KA5 (Kin.Dom.2-C-terminal) 100 TRKA RPS6KA5 (Kin.Dom.2-C-terminal) 100 TRKA RPS6KA5 (Kin.Dom.2-C-terminal) 86 TRKD	100
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RIPK5 77 TLK1 ROCK1 92 TLK2 ROCK2 100 TNIK ROS1 100 TNK1 RPS6KA4 (Kin.Dom.1-N-terminal) 93 TNK2 RPS6KA5 (Kin.Dom.1-N-terminal) 100 TRKA RPS6KA5 (Kin.Dom.1-N-terminal) 100 TRKA RPS6KA5 (Kin.Dom.1-N-terminal) 100 TRKB RSK1 (Kin.Dom.1-N-terminal) 86 TRKC	100
ROCK1 92 TLK2 ROCK2 100 TNIK ROS1 100 TNK1 RPS6KA4 (Kin.Dom.1-N-terminal) 93 TNK2 RPS6KA4 (Kin.Dom.2-C-terminal) 80 TNNI3K RPS6KA5 (Kin.Dom.1-N-terminal) 100 TRKA RPS6KA5 (Kin.Dom.2-C-terminal) 100 TRKB RSK1 (Kin.Dom.1-N-terminal) 86 TRKC	100
ROCK2 100 TNIK ROS1 100 TNK1 RPS6KA4 (Kin.Dom.1-N-terminal) 93 TNK2 RPS6KA5 (Kin.Dom.2-C-terminal) 80 TNNI3K RPS6KA5 (Kin.Dom.1-N-terminal) 100 TRKA RPS6KA5 (Kin.Dom.2-C-terminal) 100 TRKB RSK1 (Kin.Dom.1-N-terminal) 86 TRKC	100
ROS1 100 TNK1 RPS6KA4 (Kin.Dom.1-N-terminal) 93 TNK2 RPS6KA4 (Kin.Dom.2-C-terminal) 80 TNNI3K RPS6KA5 (Kin.Dom.1-N-terminal) 100 TRKA RPS6KA5 (Kin.Dom.1-N-terminal) 100 TRKB RSK1 (Kin.Dom.1-N-terminal) 86 TRKC	92
RPS6KA4 (Kin.Dom.1-N-terminal) 93 TNK2 RPS6KA4 (Kin.Dom.2-C-terminal) 80 TNNI3K RPS6KA5 (Kin.Dom.1-N-terminal) 100 TRKA RPS6KA5 (Kin.Dom.1-N-terminal) 100 TRKB RSK1 (Kin.Dom.1-N-terminal) 86 TRKC	100
RPS6KA4 (Kin.Dom.2-C-terminal)80TNNI3KRPS6KA5 (Kin.Dom.1-N-terminal)100TRKARPS6KA5 (Kin.Dom.2-C-terminal)100TRKBRSK1 (Kin.Dom.1-N-terminal)86TRKC	100
RPS6KA5 (Kin.Dom.1-N-terminal)100TRKARPS6KA5 (Kin.Dom.2-C-terminal)100TRKBRSK1 (Kin.Dom.1-N-terminal)86TRKC	100
RPS6KA5 (Kin.Dom.2-C-terminal)100TRKBRSK1 (Kin.Dom.1-N-terminal)86TRKC	100
RSK1 (Kin.Dom.1-N-terminal) 86 TRKC	82
	100
RSK1 (Kin.Dom.2-C-terminal) 100 TRPM6	100
RSK2 (Kin.Dom.1-N-terminal) 83 TSSK1B	100
RSK3 (Kin.Dom.1-N-terminal) 100 TTK	70
RSK3 (Kin.Dom.2-C-terminal) 93 TXK	100
RSK4 (Kin.Dom.I-N-terminal) 98 TYK2 (JH1domain-catalytic)	97
RSK4 (Kin.Dom.2-C-terminal) 88 TYK2 (JH2domain-pseudokinase)	84
S6K1 100 TYRO3	100
SBK1 100 ULK1	71
100 ULK2	96
SGK3 100 ULK3	100
SIK 88 VEGER2	100
SIK2 89 VRK2	33
SLK 100 WEEL	100
NARK 100 WEE2	88
SNRK 100 YANKI	100
SRC 96 YANK2	100
SRMS 100 VANK3	94
SRPK1 100 YES	100
SRPK2 66 YSK1	75
SRPK3 76 VSK4	100
5TK16 100 74K	100
STK33 90 ZAP70	59
STK35 100	

AUTOPHAGY STUDIES



Supplementary Figure 1. Aftin-5 is not involved in autophagy. A) Immunofluorescence of N2A-A β PP695 cells treated with the indicated drug for 18 h (A-5: Aftin-5; Rapa: Rapamycin). Cells were stained for LC3 (green), DNA is visualized with DAPI (blue). Bar: 10 μ m. B) Quantification of LC3 foci present per cell following treatment with 0.6 μ M Rapamycin, 100 μ M Aftin-5 or 100 μ M MRT2-309 (2), an inactive analogue of Aftin-5, or DMSO vehicle (control). When cells were treated with two drugs, rapamycin was added two hours after Aftin-5 or MRT2-309.

MITOCHONDRIAL STUDIES

Material and methods - Results

Assessment of mitochondrial swelling in cultured cells

N2a-AβPP695 cells were treated with Aftin-5 (25, 50, and 100 μM) in presence or absence of cyclosporine A (10 μM; PTP inhibitor) for 6 and 24 h. Then cells were fixed with 2% glutaraldehyde in 0.1 M Na-cacodylate buffer, pH 7.2 for 3 h at 4°C. After 2 washes with 0.2 M sucrose in 0.1 M Na-cacodylate buffer, pH 7.2, the specimens were post-fixed with 1% osmium tetroxide containing 1.5% potassium cyanoferrate, dehydrated in gradual ethanol (30–100%) and embedded in Epon. 70 nm thin sections were collected onto 200 mesh cupper grids, counterstained with uranyl acetate and lead citrate before examination with a Zeiss EM 902 transmission electron microscope at 80 Kvolt (MIMA2, Electron Microscopy Platform-GPL, Jouy-en-Josas).

Assessment of $\Delta \Psi_m$ loss in cultured cells

N2a-A β PP695 cells were treated with Aftin-5 for 6 and 24 h before measure of mitochondrial

transmembrane potential $(\Delta \Psi_m)$ which was monitored by flow cytometry (FACSCalibur; BD Biosciences) using 10 nM Dioc-6 dye. N2a-A β PP695 cells treated with 1 μ M staurosporine were used as a positive control for 100% $\Delta \Psi_m$ loss.

Determination of mitochondrial cytochrome c release in cultured cells

N2a-A β PP695 cells were treated with Aftin-5 for 6 and 24 h, then permeabilized by digitonin, fixed and labeled with anti-cytochrome *c* antibody (BD Pharmingen) and secondary antibody (Alexa Fluor 488) for detection of cytochrome *c* by flow cytometry (FAC-SCalibur; BD Biosciences). Cells treated with 1 μ M staurosporine were used as a positive control for 100% cytochrome *c* release.

Measurement of oxygen consumption in cultured cells

N2a-AβPP695 cells were treated with Aftin-5 for 6 and 24 h before measure of cellular respiration. Cells were incubated in presence of the oxygen-sensitive fluorescent dye MitoXpress (Luxcel, Cork, Ireland) in 96-well plates. Oxygen consumption was measured in real-time by spectrofluorimetry (Tecan Infinite 200; $\lambda_{Excitation}$ 380 nm; $\lambda_{Emission}$ 650 nm). Treatment with rotenone 2 μ M was used as positive control for 100% inhibition of oxygen consumption.

Purification of N2a-A βPP695 mitochondria

Mitochondria were isolated from N2a-AβPP695 cell line as previously described [3]. To ensure quality of mitochondrial preparations, samples were subjected to various assays for integrity and functionality as described in Lecoeur et al. [4] and Buron et al. [3].

Assessment of swelling and $\Delta \Psi_m$ loss in isolated mitochondria

Mitochondrial swelling and $\Delta \Psi_m$ were evaluated as described previously [5] in the presence of succinate and rotenone. Calcium (CaCl₂; 50 µM) and mCICCP (50 µM) were used as the 100% baseline for swelling and loss of $\Delta \Psi_m$, respectively.

Determination of cytochrome c release in isolated mitochondria

Cytochrome *c* release was evaluated as previously described [5] using an ELISA kit (R&D Systems, France). Treatment with 20 μ g/mL Alamethicin was used as the 100% baseline.

Measurement of oxygen consumption in isolated mitochondria

Oxygen consumption was monitored as previously described [5] using the oxygen-sensitive phosphorescent dye MitoXpress (Luxcel, Cork, Ireland). Treatment with rotenone 2 μ M and Oligomycin A 1 μ M were used as positive control for 100% inhibition of oxygen consumption by complex I and complex II respectively. Untreated mitochondria were the 0% activation/inhibition of oxygen consumption. Results are presented in Supplementary Table 2.

RESULTS

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SUDD	ementary	Table	- 2

Effects of Aftin-5 on mitochondria from N2a-A β PP695 cells. Aftin-5 was tested for its ability to induce swelling, loss of $\Delta \Psi_m$, cytochrome c release, and inhibition of mitochondrial respiration. Assays were performed in both whole cells (6 and 24 h-treatment; concentration range from 1.56 to 100 μ M) and isolated mitochondria (concentration range from 3.13 to 200 μ M). The results shown are for 100 μ M on cultured cells (24 h treatment) and for 200 μ M on isolated mitochondria and are means of 3 independent experiments

	Mitochondrial effects of Aftin-5				
	Swelling	Swelling $\Delta \Psi_m$ lossCyto c O_2 consumption		nption	
Culured cells (24 h-100 µM	No effect	No effect	No effect	34% inhibition	
Isolated mitochondria (200 µM)	No effect	No effect	No effect	31% activation Complex I	No effect Complex II

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