

## Supplementary Information

### **Decreased *Npas4* and *Arc* mRNA levels in the hippocampus of aged memory-impaired wild type but not memory preserved 11 $\beta$ -HSD1 deficient mice**

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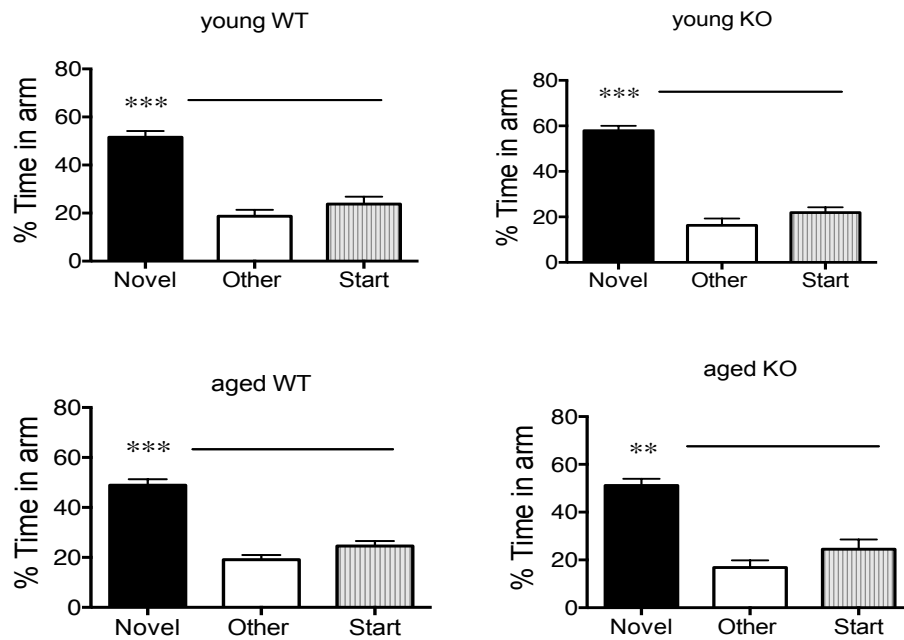
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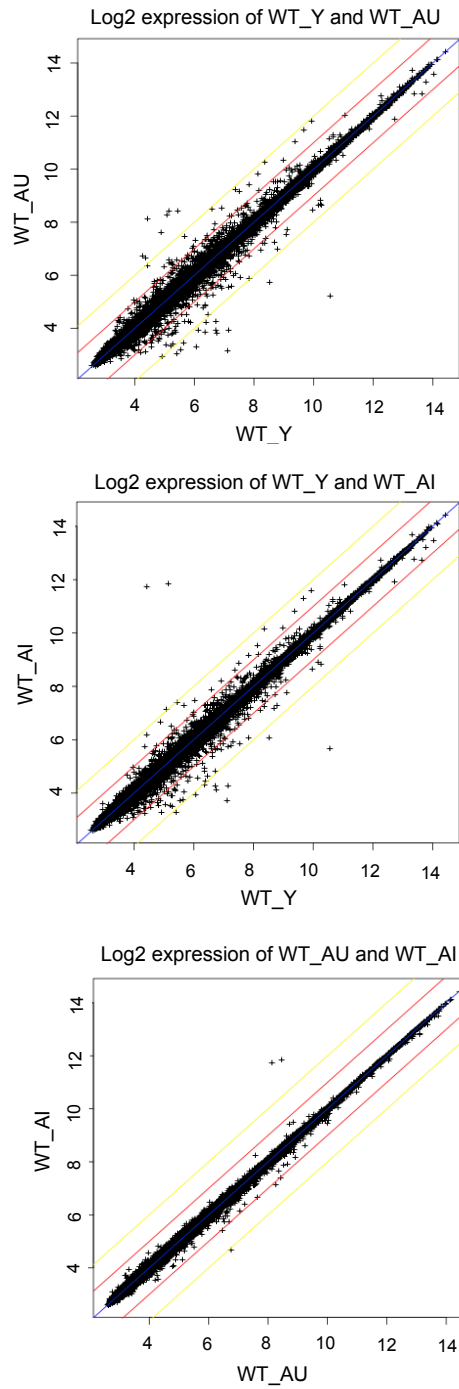
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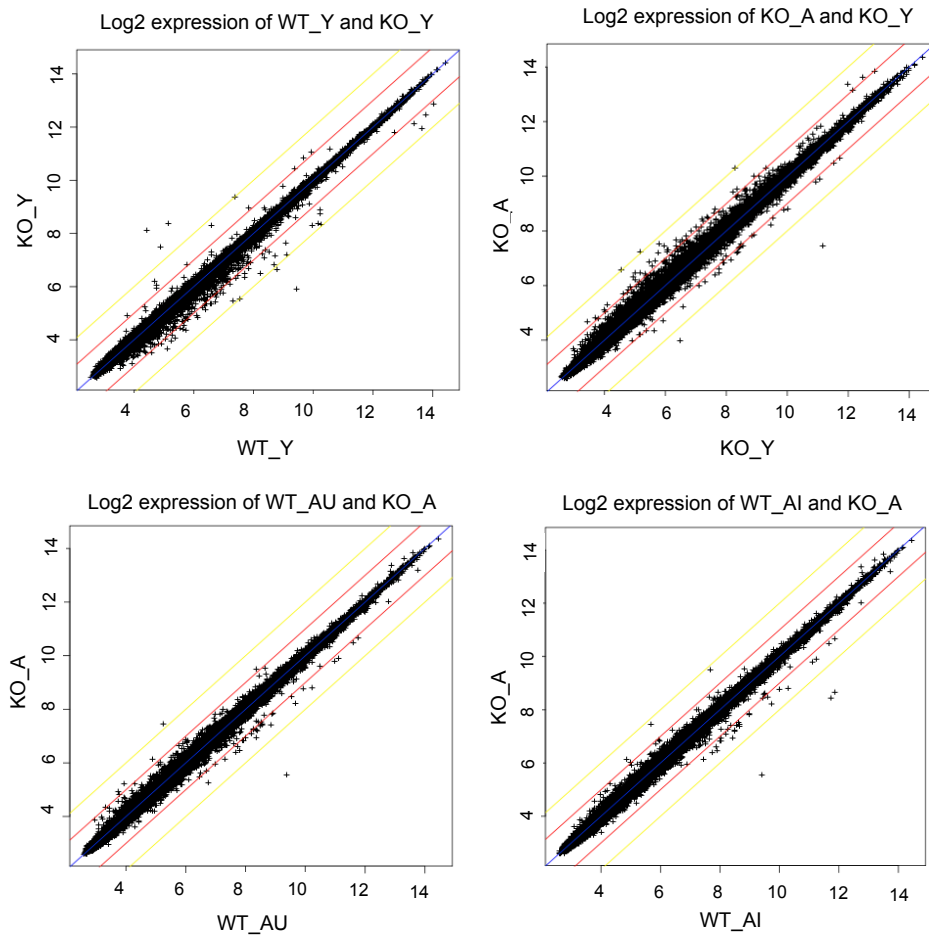
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**Figure S1. Initial assessment of young and aged wild-type and 11 $\beta$ -HSD1-deficient mice in the Y-maze following a 1 min inter-trial interval (ITI).** All young (6 months, n=9/genotype) and aged (24 months, wild-type (WT) (n=13); *Hsd11b1*<sup>-/-</sup> (KO) (n=8)) mice showed a preference for the novel arm after a 1min ITI. \*\*\* P<0.001; \*\*P<0.01 compared to other and start arms. Data shown are mean  $\pm$  SEM.



**Figure S2. Comparison of scatter plots of log intensity values (four replicates combined) for young and aged wild-type mice main comparisons.** WT\_AI (aged wild-type memory-impaired); WT\_AU (aged wild-type memory-unimpaired); WT\_Y (young wild-type). Blue line shows no change, red lines equal a 2-fold change, yellow lines are 4-fold



**Figure S3. Comparison of scatter plots of log intensity values (four replicates combined) for young and aged wild-type and 11 $\beta$ -HSD1-deficient mice main comparisons.**

WT\_AI (aged wild-type memory-impaired); WT\_AU (aged wild-type memory-unimpaired); WT\_Y (young wild-type); KO\_Y (young *Hsd11b1*<sup>-/-</sup>); KO\_A (aged *Hsd11b1*<sup>-/-</sup>)

Blue line shows no change, red lines equal a 2-fold change, yellow lines are 4-fold

## Supplementary Table S1

Hippocampal genes up-regulated with ageing in wild-type and 11 $\beta$ -HSD1-deficient mice but not affected by genotype. Gene expression level fold changes shown are from comparisons of aged (24 months) wild-type (WT\_A) mice (combined WT\_AU and WT\_AI mice) with corresponding young (6 months) wild-type (WT) or *Hsd11b1*<sup>-/-</sup> (KO) mice. \*, P<0.05, \*\*P<0.01, RP scores

Affymetrix ID	Gene symbol	Gene title	Fold change WT	Fold change KO
1426508_at	<i>Gfap</i>	glial fibrillary acidic protein	1.6 *	2 **
1449401_at	<i>C1qc</i>	complement component 1, q subcomponent, C chain	1.6 *	1.7 **
1417063_at	<i>C1qb</i>	complement component 1, q subcomponent, beta polypeptide	1.5 *	1.6 **
1419100_at	<i>Serpina3n</i>	serine (or cysteine) peptidase inhibitor, clade A, member 3N	2.0 *	1.8 *
1449254_at	<i>Spp1</i>	secreted phosphoprotein 1	1.7 *	1.6 *
1427371_at	<i>Abca8a</i>	ATP-binding cassette, sub-family A (ABC1), member 8a	1.6 **	1.8 **
1452428_a_at	<i>B2m</i>	beta-2 microglobulin	1.5 *	1.6 *
1418204_s_at	<i>Aif1</i>	allograft inflammatory factor 1	1.5 *	1.7 **
1435477_s_at	<i>Fcgr2b</i>	Fc receptor, IgG, low affinity IIb	1.9 *	2.0 **
1444564_at	<i>Apod</i>	apolipoprotein D	2.0 *	2.0 **
1450641_at	<i>Vim</i>	Vimentin	1.6 *	1.8 **
1422640_at	<i>Pcdhb9</i>	protocadherin beta 9	2.0 **	1.9 **
1454268_a_at	<i>Cyba</i>	cytochrome b-245, alpha polypeptide	1.6 *	1.8 **
1448620_at	<i>Fcgr3</i>	Fc receptor, IgG, low affinity III	1.5 *	1.7 **
1460330_at	<i>Anxa3</i>	annexin A3	1.4 *	1.6 **
1421792_s_at	<i>Trem2</i>	triggering receptor expressed on myeloid cells 2	1.5 **	1.5 *
1418580_at	<i>Rtp4</i>	receptor transporter protein 4	2.0 *	2.0 **
1443745_s_at	<i>Dmp1</i>	dentin matrix protein 1	1.6 *	1.9 **
1423547_at	<i>Lyz2</i>	lysozyme 2	2.9 **	2.4 *
1427076_at	<i>Mpeg1</i>	macrophage expressed gene 1	1.6 *	2.0 **
1450792_at	<i>Tyrobp</i>	TYRO protein tyrosine kinase binding protein	1.6 *	1.8 **
1434342_at	<i>S100b</i>	S100 protein, beta polypeptide, neural	1.2 *	1.7 **