

## **Supporting information**

### ***Hansenula polymorpha* Aat2p is targeted to peroxisomes via a novel Pex20p dependent pathway**

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## **Supporting materials and methods**

### *Protein purification and preparation of antibodies*

*H. polymorpha* Aat2p with a cleavable His<sub>6</sub>-GST tag was produced in *E. coli* BL21 (DE3) RIL. Cells were grown at 37°C to an OD<sub>600</sub> of 0.6 in Terrific Broth (TB) medium, transferred to 21°C and grown until an OD<sub>600</sub> of 1.5. Gene expression was induced with 0.05mM IPTG (Invitrogen) for 16 h and cells were harvested by centrifugation. Cell pellets were resuspended in lysis buffer (50mM Tris-HCl pH 7.5, 150mM NaCl, 1% glycerol, 1mM DTT, 1mg/ml lysozyme, 10µg/ml DNase) and passed two times through a French press. Cell debris was removed by centrifugation and lysates were loaded onto glutathione sepharose-4B resin (GE Healthcare) pre-equilibrated with lysis buffer. The resin was extensively washed with lysis buffer and His<sub>6</sub>-GST tagged proteins were eluted using lysis buffer containing 20mM reduced glutathione. The GST tag was cleaved from the protein using TEV protease and samples were passed over a Ni-NTA column. The concentrated protein sample was subjected to gel filtration using a Superdex 200 (16/60) column (GE Healthcare) equilibrated with 25mM Tris, 150mM NaCl, 1mM 2-mercaptoethanol, pH 7.5. The presence of purified

Aat2p was confirmed using SDS-PAGE and protein samples were sent for antibody production (Eurogentec). The properties of the resulting anti-Aat2p antibodies are shown in Figure S1.

*Construction of strains containing N-terminally tagged GFP-Aat2*

The N-terminal GFP fusion of Aat2p produced from its endogenous promoter was obtained by amplifying a 0.7 kb fragment of GFP along with the restriction sites HindIII and BamHI using pHIPZ-Pex13mGFP as template and the primers ANN PR46 and ANN PR47. Similarly, a fragment of 1.3 kb corresponding to *AAT2* was amplified incorporating BamHI and XbaI restriction sites from genomic DNA using primers ANN PR48 and ANN PR49. HindIII-BamHI digested GFP was ligated into HindIII-BamHI digested pUC19 and BamHI-XbaI digested *AAT2* was ligated with BamHI-XbaI digested pUC19. Following this, GFP was excised using HindIII and BamHI, *AAT2* was excised using BamHI and XbaI and these fragments were used in a three-way ligation with HindIII-XbaI digested pHIPN5 GFP-SKL [1] to yield pHIPN5-mGFPAat2 (pANN017). To produce the fusion protein from the endogenous promoter, the GFP-*AAT2* fragment was excised with HindIII and NsiI and ligated with HindIII-NsiI digested pANN015, resulting in the plasmid pHIPNP<sub>*AAT2*</sub>-mGFPAat2 (pANN014). StuI was used for linearization before integration into *H. polymorpha yku80* and *aat2* cells.

The plasmid pHIPX7DsRed-SKL was linearized with DraI prior to transformation into WT+GFP-AAT2 and *aat2*+GFP-AAT2 cells.

**Table S1: *H. polymorpha* strains used in this study.**

Strains	Characteristics	Reference
<i>yku80</i> (WT)	NCYC495, <i>leu1.1</i> YKU80:: <i>URA3</i>	[2]
WT+AAT2-GFP	YKU80:: <i>URA3</i> + pANN009	This study
<i>pex1</i>	NCYC495, <i>leu1.1</i> PEX1:: <i>URA3</i>	[3]
<i>pex1.atg1</i>	NCYC495, <i>leu1.1</i> PEX1:: <i>URA3</i> , ATG1:: <i>HPH</i>	This study
<i>pex2</i>	NCYC495, <i>leu1.1</i> PEX2:: <i>URA3</i>	[4]
<i>pex3</i>	NCYC495, <i>leu1.1</i> PEX3:: <i>URA3</i>	[5]
<i>pex5</i> ( <i>LEU2</i> marker, used for construction of <i>pex5.pex7</i> )	NCYC495, <i>ura3</i> PEX5:: <i>LEU2</i>	[6]
<i>pex5.pex7</i>	Segregant of cross between PEX5:: <i>LEU2</i> and PEX7:: <i>URA3</i>	This study
<i>pex13</i>	NCYC495, <i>leu1.1</i> PEX13:: <i>URA3</i>	[4]
<i>pex19</i>	NCYC495, <i>leu1.1</i> PEX19:: <i>URA3</i>	[7]
<i>pex20</i>	NCYC495, <i>leu1.1</i> PEX20:: <i>URA3</i>	[8]
WT+AAT2-GFP+Pex14-mCherry	YKU80:: <i>URA3</i> + pANN009+pHIPN-Pex14-mCherry	This study
<i>pex1.atg1</i> +AAT2-GFP+Pex14-mCherry	PEX1:: <i>URA3</i> ,ATG1:: <i>HPH</i> + pANN009+pHIPN-Pex14-mCherry	This study
<i>pex2</i> +AAT2-GFP+Pex14-mCherry	PEX2:: <i>URA3</i> + pANN009+pHIPH-Pex14-mCherry	This study
<i>pex5.pex7</i> +AAT2-GFP+Pex14-mCherry	PEX5:: <i>LEU2</i> ,PEX7:: <i>URA3</i> + pANN009+pHIPN-Pex14-mCherry	This study
<i>pex13</i> +AAT2-GFP+Pex14-mCherry	PEX13:: <i>URA3</i> + pANN009+pHIPH-Pex14-mCherry	This study
<i>pex20</i> +AAT2-GFP+Pex14-mCherry	PEX20:: <i>URA3</i> + pANN009+pHIPN-Pex14-mCherry	This study
<i>aat2</i>	AAT2:: <i>HPH</i>	This study
WT+GFP-AAT2	YKU80:: <i>URA3</i> + pANN014	This study
WT+GFP-AAT2+DsRed-SKL	YKU80:: <i>URA3</i> + pANN014+pHIPX7-DsRed-SKL	This study
<i>aat2</i> +GFP-AAT2	AAT2:: <i>HPH</i> + pANN014	This study
<i>aat2</i> +GFP-AAT2+DsRed-SKL	AAT2:: <i>HPH</i> +pANN014+pHIPX7-DsRed-SKL	This study

**Table S2: Plasmids used in this study.**

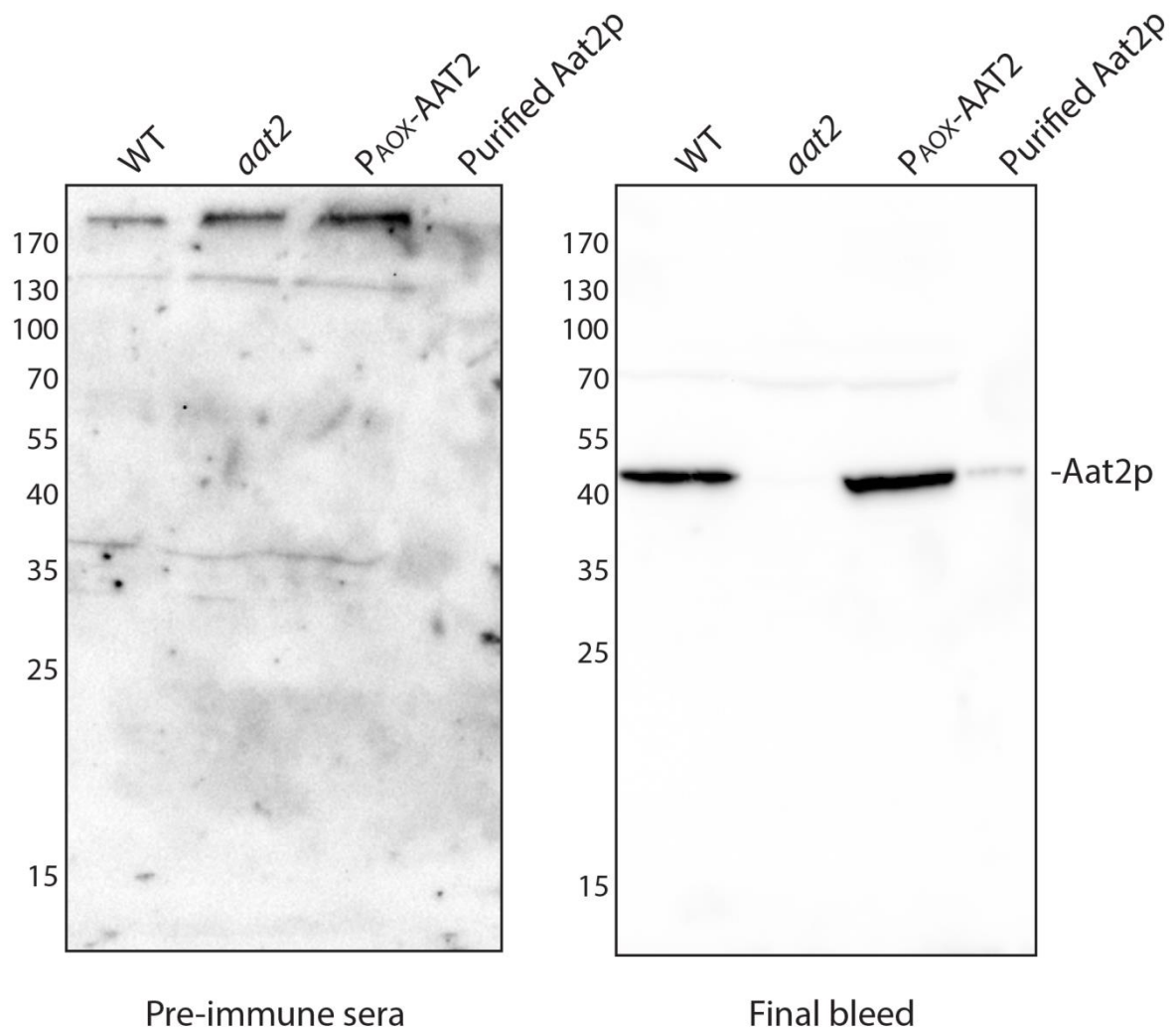
Plasmid	Description	Reference
pAG32	<i>S. cerevisiae</i> vector containing hygromycin B resistance cassette (Hph <sup>R</sup> ), Amp <sup>R</sup>	[9]
pHIPH4	pHIP vector bearing the <i>AOX</i> promoter, Amp <sup>R</sup> , Hph <sup>R</sup>	This study

pHIPZ4	pHIP vector containing the <i>AOX</i> promoter, Amp <sup>R</sup> , Zeo <sup>R</sup>	[10]
pHIPN4	Plasmid bearing the <i>AOX</i> promoter, Amp <sup>R</sup> , Nat <sup>R</sup>	[11]
pANN016	<i>AAT2</i> expressed from the <i>AOX</i> promoter, contains Amp <sup>R</sup> , Nat <sup>R</sup>	This study
pHIPZ-Pex13mGFP	<i>PEX13</i> expressed from the endogenous promoter with C-terminal GFP fusion, contains Amp <sup>R</sup> , Zeo <sup>R</sup>	[12]
pANN009	<i>AAT2</i> expressed from the endogenous promoter with C-terminal GFP fusion, contains Amp <sup>R</sup> , Zeo <sup>R</sup>	This study
pHIPN-Pex14-mCherry	<i>PEX14</i> expressed from the endogenous promoter with C-terminal mCherry fusion, contains Amp <sup>R</sup> , Nat <sup>R</sup>	[12]
pHIPH-Pex14-mCherry	<i>PEX14</i> expressed from the endogenous promoter with C-terminal mCherry fusion, contains Amp <sup>R</sup> , Hph <sup>R</sup>	This study
pANN015	<i>AAT2</i> expressed from the endogenous promoter, contains Amp <sup>R</sup> , Nat <sup>R</sup>	This study
pDONR P4-P1R	Multisite Gateway vector; Kan <sup>R</sup> , Cm <sup>R</sup>	Invitrogen
pDONR P2R-P3	Multisite Gateway vector; Kan <sup>R</sup> , Cm <sup>R</sup>	Invitrogen
pENTR <i>ATG1</i> 5'	pDONR P4-P1R with 5' flanking region of <i>ATG1</i> ; Kan <sup>R</sup>	This study
pENTR <i>ATG1</i> 3'	pDONR P2R-P3 with 3' flanking region of <i>ATG1</i> ; Kan <sup>R</sup>	This study
pDEST-R4-R3	Multisite Gateway donor vector; Amp <sup>R</sup> , Cm <sup>R</sup>	Invitrogen
pENTR221-hph	pDONR 221 with <i>HPH</i> ; Hph <sup>R</sup> , Kan <sup>R</sup>	[2]
pARM011	Plasmid bearing the <i>ATG1</i> deletion cassette, contains Amp <sup>R</sup> , Hph <sup>R</sup>	This study
pETM30 <i>AAT2</i>	His <sub>6</sub> GST- <i>Aat2</i> , for expression in <i>E. coli</i> ; Kan <sup>R</sup>	This study
pHIPN5GFP-SKL	GFP-SKL expressed from the <i>AMO</i> promoter, contains Amp <sup>R</sup> , Nat <sup>R</sup>	[1]
pANN017	<i>AAT2</i> expressed from the <i>AMO</i> promoter with N-terminal GFP fusion, contains Amp <sup>R</sup> , Nat <sup>R</sup>	This study
pANN014	<i>AAT2</i> expressed from endogenous promoter, with N-terminal GFP fusion, contains Amp <sup>R</sup> , Nat <sup>R</sup>	This study
pHIPX7DsRed-SKL	DsRed-SKL expressed from the constitutive <i>TEF</i> promoter, contains Kan <sup>R</sup> , LEU2	[13]

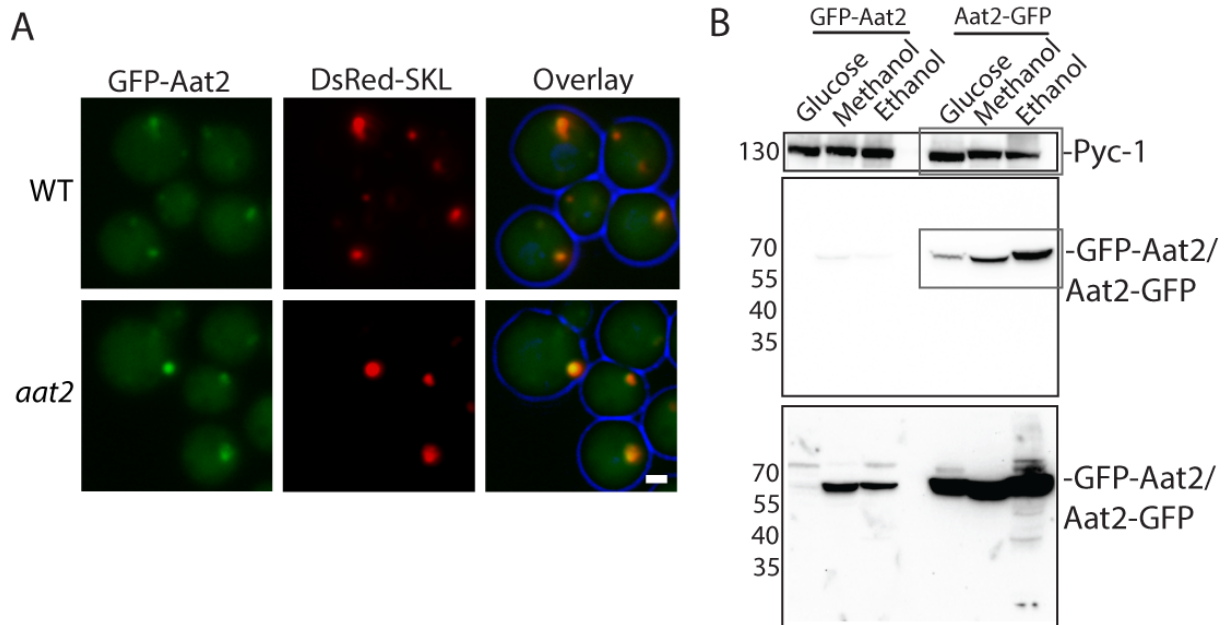
**Table S3: Oligonucleotides used in this study.**

Primer	Sequence
ANN PR15	GATTTTGCCCTCTGTCAGGCTCGCCGAGAACCTGTTGCAGAACTC CAAGGCCACACACCATAGCTTCAA
ANN PR16	TCGTCCTCCCCTGTTTCATCAGCTCGGGCGTGTTGAGGATCAGA GACACCGTTTTGACACTGGATGGC

ANN PR27	CACGTAAAGCTTACAGACTCGTGTCTATCCAG
ANN PR28	GATACGAGATCTCACACTTCTCACACCTCGTC
ANN PR35	GCGCAAGCTTATGACAAGATCCTTCAGCATCGAGAACATCC
ANN PR51	CGCTCTAGACTACACACTTCTCACACCTCGTCAATACAT
ARM PR 16	GGGGACAGCTTTCTTGTACAAAGTGGCCGCCACAAATGGTGAAG TCGATC
ARM PR 17	GGGGACAACCTTTGTATAATAAAGTTGCATCGAGCTTCTCGTTGCC CGTGAC
ARM PR 18	GGGGACAACCTTTGTATAGAAAAGTTGGGCTGGAGAACGCGGCAG ATCC
ARM PR 19	GGGGACTGCTTTTTTGTACAACTTGGGGAGGGGAAGGGTACCT CTC
ARM PR 20	ACAGGTCGTTGGTGACTTTAC
ARM PR 21	CTTCTCGTTGCCCGTGACC
ARM PR 62	AGGCCGTTGCGTTATGATAG
ARM PR 63	GGACAATGGCCGCATAACAG
PRARM001	ATAGCGGCCGCTTGCAGGAAGTCGACGAAAT
PRARM002	CGGAAGCTTTTACTTGTACAGCTCGTCCA
AAT2 Ab_F	CATGCCATGGCAATGACAAGATCCTTCAGCATCG
AAT2 Ab_R	CCCAAGCTTCTACACACTTCTCACACCTCG
ANN PR83	AAGCGGCCGCGATTTTCGGGTCCAGAGTGT
ANN PR84	GCAAGCTTGGGGGAGAAATGGGAGGAAG
ANN PR 46	CGCAAGCTTATGGTGAGCAAGGGCGAG
ANN PR 47	GCGCGGATCCCTTGTACAGCTCGTCCATG
ANN PR48	GCGCGGATCCACAAGATCCTTCAGCATCGAG
ANN PR49	GCGCTCTAGACTACACACTTCTCACACC



**Figure S1: Specificity of *HpAat2p* antibodies.** Western blot analysis of cell lysates from methanol grown WT cells, *aat2* cells and WT cells containing *AAT2* under control of the strong alcohol oxidase promoter (*P<sub>AOX</sub>-AAT2*), together with purified Aat2p from *E.coli*, probed with pre-immune serum (left panel) or serum from a rabbit immunogenized with purified Aat2p (right panel). Blots were exposed for 20 sec.



**Figure S2: N-terminally tagged GFP-Aat2 partially localises to peroxisomes. (A)** Co-localisation analysis of GFP-Aat2 in ethanol-grown WT (upper panel) or *aat2* (lower panel) cells co-expressing the peroxisomal matrix marker DsRed-SKL. Scale bar represents 1µm. **(B)** Cell lysates of WT cells grown on different carbon sources expressing Aat2p tagged with GFP either at the N- or C-terminus were subjected to SDS-PAGE and immunoblotting using antibodies directed against GFP or Pyc-1 (loading control). The two blots represent different exposures: top panel, 10 sec. exposure, bottom panel, 1000 sec. exposure. Part of this blot (boxed) is shown in Figure 2A.

## Supporting references

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