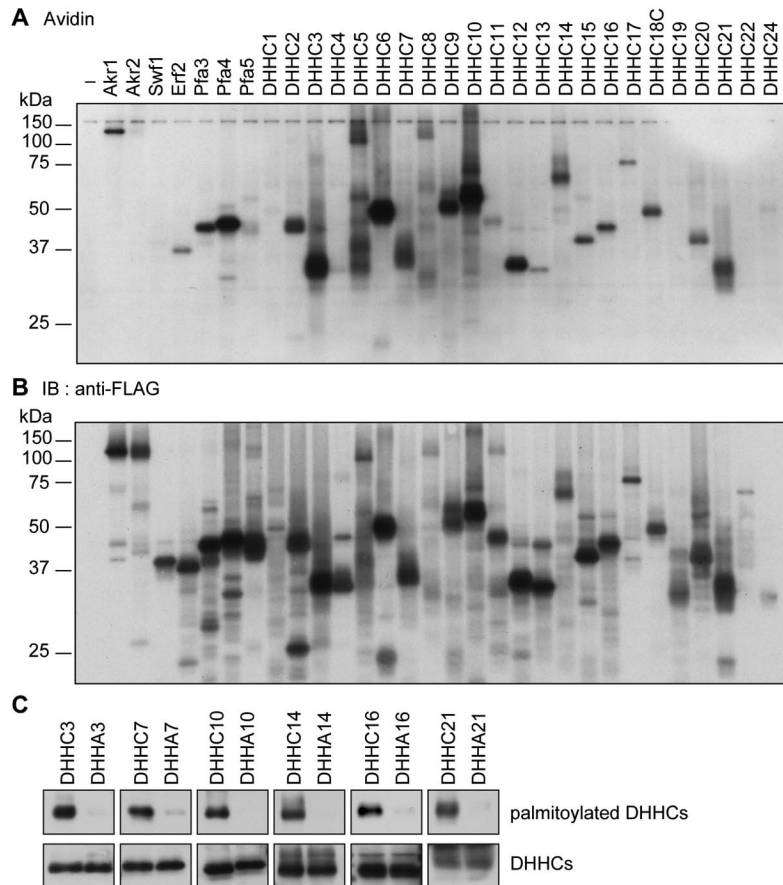


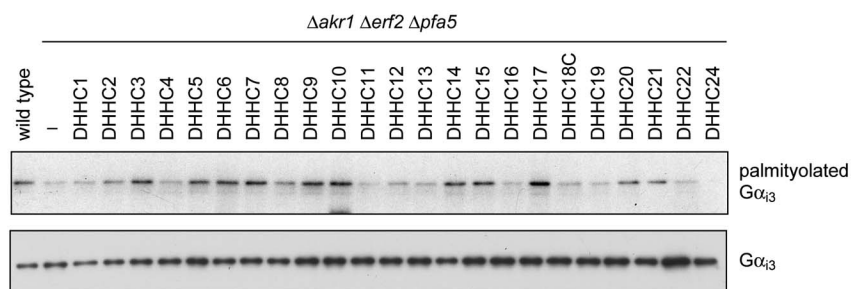
## **Supplemental Materials**

### **Analysis of substrate specificity of human DHHC protein acyltransferases using a yeast expression system**

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**Fig. S1.** Twenty human DHHC proteins form acyl intermediates in yeast. (A and B) SEY6210 (wild type) cells were transfected with the pAKNF314 (vector), pAKNF-AKR1 (*3xFLAG-AKR1*), pAKNF-AKR2 (*3xFLAG-AKR2*), pAKNF-SWF1 (*3xFLAG-SWF1*), pAKNF-ERF2 (*3xFLAG-ERF2*), pAKNF-PFA3 (*3xFLAG-PFA3*), pAKNF-PFA4 (*3xFLAG-PFA4*), pAKNF-PFA5 (*3xFLAG-PFA5*), pAKNF-DHHC<sub>x</sub> (*3xFLAG-DHHC<sub>x</sub>* with x representing the DHHC number; x=1-17, 19-22, and 24), or pLECF-DHHC18 (*DHHC18-3xFLAG*) plasmid. Total cell lysates were treated with *N*-ethylmaleimide (NEM) to block free Cys residues. After removal of palmitic acid from the palmitoylated Cys residues by incubation with hydroxylamine, the exposed free Cys residues were modified with biotin using Biotin-HPDP. The samples were then subjected to immunoprecipitation with anti-FLAG antibodies, followed by detection with streptavidin-HRP (A) or with immunoblotting using anti-FLAG antibody (B). (C) SEY6210 cells were transfected with pAKNF-DHHC<sub>x</sub> or pAKNF-DHHAx (x represents the DHHC number; x=3, 7, 10, 14, 16, and 21) plasmid. Total cell lysates were prepared and treated sequentially with NEM, hydroxylamine, and Biotin-HPDP. The biotinylated samples were precipitated with immobilized avidin beads and detected by immunoblotting with anti-FLAG antibody (upper panel). The DHHC proteins in the total lysates were also detected using immunoblotting with anti-FLAG antibody (lower panel).



**Fig. S2.** Thirteen human DHHC proteins palmitoylate  $G\alpha_{i3}$ . The KAY251 ( $\Delta akr1 \Delta erf2 \Delta pfa5$ ) cells harboring the pYU- $G\alpha_{i3}$  ( $G\alpha_{i3}$ -9xMyc) plasmid were transfected with the pAKNF314 (vector), pAKNF-DHHCx ( $3xFLAG$ -DHHCx with x representing the DHHC number; x=1-17, 19-22, and 24), or pLECF-DHHC18 ( $DHHC18$ -3xFLAG) plasmid. Total cell lysates were prepared from the transfected cells, as well as wild type (SEY6210) cells bearing the pYU- $G\alpha_{i3}$  and pAKNF314 plasmids, and treated sequentially with NEM, hydroxylamine, and Biotin-HPDP. The DHHC proteins were immunoprecipitated with anti-Myc antibodies, and the precipitates were detected by Ultra Sensitive ABC Peroxidase Staining Kit (upper panel) or immunoblotting with anti-Myc antibody (lower panel).

**Table S1. Human and mouse *DHHC* genes**

DHHC <sup>a</sup>		ZDHHC <sup>b</sup>		Other Name	Accession Number	
Human	Mouse	Human	Mouse		Human	Mouse
<i>DHHC1</i>	<i>DHHC1</i>	<i>ZDHHC1</i>	<i>Zdhhc1</i>		NM_013304	BC026570
<i>DHHC2</i>	<i>DHHC2</i>	<i>ZDHHC2</i>	<i>Zdhhc2</i>	<i>REAM</i>	BC050272	NM_178395
<i>DHHC3</i>	<i>DHHC3</i>	<i>ZDHHC3</i>	<i>Zdhhc3</i>	<i>GODZ</i>	NM_016598	NM_026917
<i>DHHC4</i>	<i>DHHC4</i>	<i>ZDHHC4</i>	<i>Zdhhc4</i>		NM_018106	NM_028379
<i>DHHC5</i>	<i>DHHC5</i>	<i>ZDHHC5</i>	<i>Zdhhc5</i>		NM_015457	NM_144887
<i>DHHC6</i>	<i>DHHC6</i>	<i>ZDHHC6</i>	<i>Zdhhc6</i>		BC007213	NM_025883
<i>DHHC7</i>	<i>DHHC7</i>	<i>ZDHHC7</i>	<i>Zdhhc7</i>	<i>SERZ-β</i>	NM_017740	NM_133967
<i>DHHC8</i>	<i>DHHC8</i>	<i>ZDHHC8</i>	<i>Zdhhc8</i>		NM_013373	AY668947
<i>DHHC9</i>	<i>DHHC9</i>	<i>ZDHHC9</i>	<i>Zdhhc9</i>		BC012826	AK032233
<i>DHHC10</i>	<i>DHHC10</i>	<i>ZDHHC11</i>	<i>Zdhhc11</i>		NM_024786	AY668948
<i>DHHC11</i>	<i>DHHC11</i>	<i>ZDHHC23</i>	<i>Zdhhc23</i>	<i>NIDD</i>	NM_173570	AY668949
<i>DHHC12</i>	<i>DHHC12</i>	<i>ZDHHC12</i>	<i>Zdhhc12</i>		NM_032799	BC021432
<i>DHHC13</i>	<i>DHHC13</i>	<i>ZDHHC24</i>	<i>Zdhhc24</i>		NM_207340	BC071194
<i>DHHC14</i>	<i>DHHC14</i>	<i>ZDHHC14</i>	<i>Zdhhc14</i>		NM_024630	BC059814
<i>DHHC15</i>	<i>DHHC15</i>	<i>ZDHHC15</i>	<i>Zdhhc15</i>		BC103980	NM_175358
<i>DHHC16</i>	<i>DHHC16</i>	<i>ZDHHC16</i>	<i>Zdhhc16</i>	<i>APH2</i>	NM_032327	XM_129300
<i>DHHC17</i>	<i>DHHC17</i>	<i>ZDHHC17</i>	<i>Zdhhc17</i>	<i>HIP14, AKR1</i>	NM_015336	NM_172554
<i>DHHC18</i>	<i>DHHC18</i>	<i>ZDHHC18</i>	<i>Zdhhc18</i>		NM_032283	AY668950
<i>DHHC19</i>	<i>DHHC19</i>	<i>ZDHHC19</i>	<i>Zdhhc19</i>		NM_144637	BC049761
<i>DHHC20</i>	<i>DHHC20</i>	<i>ZDHHC20</i>	<i>Zdhhc20</i>		NM_153251	AY668951
<i>DHHC21</i>	<i>DHHC21</i>	<i>ZDHHC21</i>	<i>Zdhhc21</i>		NM_178566	NM_026647
<i>DHHC22</i>	<i>DHHC22</i>	<i>ZDHHC13</i>	<i>Zdhhc13</i>	<i>HIP14L, AKRL1</i>	AB024495	NM_028031
– <sup>c</sup>	<i>DHHC23</i>	– <sup>c</sup>	<i>Zdhhc25</i>		–	BC049767
<i>DHHC24</i>	<i>DHHC24</i>	<i>ZDHHC22</i>	<i>Zdhhc22</i>		NM_174976	NM_001080943

<sup>a</sup>The *DHHC* nomenclature followed the previous report (Fukata *et al.*, 2004).

<sup>b</sup>The *DHHC* genes are registered in GenBank as *ZDHHC* (zinc-finger, *DHHC*-type). Note that nomenclatures of *DHHC* are different from those of *ZDHHC* for some *DHHC* genes.

<sup>c</sup>The *DHHC23* gene does not exist in human genome.

**Table S2. Yeast strains used in this study**

Strain	Genotype	References
SEY6210	<i>MAT<math>\alpha</math> leu2-3,112 ura3-52 his3-<math>\Delta</math>200 trp1-<math>\Delta</math>901 lys2-801 suc2-<math>\Delta</math>9</i>	(Robinson <i>et al.</i> , 1988)
SEY6211	<i>MAT<math>\alpha</math> leu2-3,112 ura3-52 his3-<math>\Delta</math>200 trp1-<math>\Delta</math>901 ade2-101 suc2-<math>\Delta</math>9</i>	(Robinson <i>et al.</i> , 1988)
KHY1028	SEY6210, $\Delta$ <i>yck2::YCK2-K-RASB URA3</i>	This study
KHY1031	SEY6210, $\Delta$ <i>yck2::YCK2-K-RASB</i>	This study
KHY1045	SEY6210, $\Delta$ <i>akr1::HIS3</i>	This study
KHY1046	SEY6210, $\Delta$ <i>yck2::YCK2-K-RASB <math>\Delta</math>akr1::HIS3</i>	This study
RHY59	<i>MAT<math>\alpha</math> leu2-3,112 ura3-52 his3-<math>\Delta</math>200 trp1-<math>\Delta</math>90 ade2-101 suc2-<math>\Delta</math>9 <math>\Delta</math><i>akr2::KanMX4 <math>\Delta</math>swf1::KanMX4 <math>\Delta</math>erf2::LEU2 <math>\Delta</math>pfa3::KanMX4 <math>\Delta</math>pfa4::LEU2 <math>\Delta</math>pfa5::KanMX4</i></i>	This study
IAY106	RHY59, $\Delta$ <i>agp1::pTEF-9xMYC-AGP1 URA</i>	This study
KAY39	KHY1046, $\Delta$ <i>yck1::pTEF-9xMYC-YCK1 URA3</i>	This study
KAY44	SEY6210, $\Delta$ <i>agp1::pTEF-9xMYC-AGP1 URA3</i>	This study
KAY47	SEY6210, $\Delta$ <i>gpa2::pTEF-GPA2-9xMYC natNT2 URA3</i>	This study
KAY57	KHY1046, $\Delta$ <i>erf2::KanMX4 <math>\Delta</math>gpa2::pTEF-GPA2-9xMYC natNT2 URA3</i>	This study
KAY62	SEY6210, $\Delta$ <i>yck1::pTEF-9xMYC-YCK1 URA3</i>	This study
KAY110	SEY6210, $\Delta$ <i>sso1::pTEF-9xMYC-SSO1 URA3</i>	This study
KAY113	SEY6210, $\Delta$ <i>swf1::KanMX4 <math>\Delta</math>sso1::pTEF-9xMYC-SSO1 URA3</i>	This study
KAY251	KHY1046, $\Delta$ <i>erf2::LEU2 <math>\Delta</math>pfa5::KanMX4</i>	This study

**Table S3. Primers used in this study**

Primer	Nucleotide sequence
DHHC18C-F	5'-AGAATTCGCCACCATGAAGGACTGCGAG-3' ( <i>EcoRI</i> )
DHHC18C-R	5'-AAGATCTTCCACCATGCTGGCATCAGG-3' ( <i>BglIII</i> )
DHHC24-F	5'-AAGGATCCATGCTGGCCCTGCGGCTGCTCAACG-3' ( <i>BamHI</i> )
DHHC24-R	5'-TGGATCCCTACTTATCCTGCTGCTTGGAGC-3' ( <i>BamHI</i> )
PSD95-F	5'-AGCCACCATGGACTGTCTCTGTATAGTGACAAC-3'
PSD95-R	5'-TGAATTCGAGAGTCTCTCTCGGGCTGGAACC-3'
SNAP25b-F	5'-TGGATCCATGGCCGAAGACGCAGACATGC-3' ( <i>BamHI</i> )
SNAP25b-R	5'-TTTAACCACTTCCCAGCATCTTTGTTGC-3'
Gαi3-F	5'-GCCACCATGGGCTGCACGTTGAGCGCCGAG-3'
Gαi3-R	5'-AACCCGGGCAAATAAAGCCCACATTCC-3' ( <i>SmaI</i> )
AKR1-F	5'-AGGATCCATGGTGAACGAATTAGAGAATG-3' ( <i>BamHI</i> )
AKR1-R	5'-ACCTCATGAGCTCTCCTAATAGTCC-3'
AKR2-F	5'-AAGATCTATGACCAGTATGTCAATTATTG-3' ( <i>BglIII</i> )
AKR2-R	5'-AATCACGCAAATTTTTGTTGTGCAC-3'
ERF2-F	5'-GGATCCATGGCCTTGGTCTCTAGAAGGTTCG-3' ( <i>BamHI</i> )
ERF2-R	5'-GCTGCACGTTGTCTGACATCTGAAGH-3'
SWF1-F	5'-GGATCCATGTCATGGAATCTACTATTTGTGC-3' ( <i>BamHI</i> )
SWF1-R	5'-CAGGACTAATTGCTGGTAGTATGGA-3'
PFA3-F	5'-GGATCCATGAATGACAGGCTTTCGTTGACAAG-3' ( <i>BamHI</i> )
PFA3-R	5'-TGGCTACAGTACCACAGTAGATTCT-3'
PFA4-F	5'-GGATCCATGCCAGTAAAGTTAAGGTGGCCTT-3' ( <i>BamHI</i> )
PFA4-R	5'-CGAGTGCTGGAAACCAGGGAATAGA-3'
PFA5-F	5'-GGATCCATGGCTCTATCATGGAATATTCGAA-3' ( <i>BamHI</i> )
PFA5-R	5'-GTGAAGATTGGAGACGTTGCCTCAC-3'
YCK2-F1	5'-GGAAACTACCAGAATACACAACGCC-3'
YCK2-R1	5'-CAATTGCCCGGGTTACATAATTACACACTTTGTCTTTGACTTCTTTTTCT TCTTTTTACCATCTCCTAGCTTACTGAAAAACCTTTCG-3'
YCK2-F2	5'-AATAGAAAACGGAGGAGGTTTTTTG-3'
YCK2-R2	5'-CTATTTGTCTATCCAGCTAAACGTG-3'
YCK1-F	5'-CATACAAAAATTCGCGCATTTCGCTGGCCCTTTTCCCTGCTCTCCTCTTCC CCATGGCCAAGCTTTTCAATTCATC-3'
YCK1-R	5'-TTATATTGGTGAGGTTGTTAACTGCTAGAGTGGTACTTGCTATGGGCAT GGACATAGATCTGCTAG-3'
GPA2-F1	5'-CGCCATCAAAGAAACAATTTGGAAAATACATTGAAAGACTCTGGAGT GTTACAACGTACGCTGCAGGTCGACTC-3'
GPA2-R1	5'-ATAATAAAGGGGAGAAGAGGCATGCAGTTTTGTCTCTGTTTTAGCTGT GCATTCACATCGATGAATTCGAGCTCG-3'

GPA2-F2 5'-CAGCGAGCCTTATTGTTACAGCACAAATCACGCGTATTTTCAAGCAAA  
TATCATGGCCAAGCTTTTCAATTCATC-3'

GPA2-R2 5'-CGGTCTGCGTGTGTCAGGAGTGCTGCCGTTCTTTTCTGAAGATGCGCAGAG  
ACCCATTCCACTAGAAAAGTTAG-3'

AGP1-F 5'-TGTAATCTTTATAGAAGAAGCACGCTAATATAGACAAAGATAGCTTCG  
CACAATGGCCAAGCTTTTCAATTCATC-3'

AGP1-R 5'-CTGTGGAGCTATTTTCAAGTCTTTCAGTTCGTATAGAGACTTCGACGA  
CGACATAGATCTGCTAGTGGATCCG-3'

SSO1-F1 5'-AAGAAAACCCTTTTACAATTAATAAAGGCAATTAATAAATAGAAACAA  
ATCAAATGGCCAAGCTTTTCAATTCATC-3'

SSO1-R1 5'-CCAACTCGTATGACTCTTCAAAGGGTTTCCAACTGGTACGGATTATT  
ATAACTAGATCTGCTAGTGGATCCG-3'

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### **Supplemental References**

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Robinson JS, Klionsky DJ, Banta LM, Emr SD (1988). Protein sorting in *Saccharomyces cerevisiae*: isolation of mutants defective in the delivery and processing of multiple vacuolar hydrolases. *Mol Cell Biol* 8, 4936-4948.