

# Supplemental Materials

*Molecular Biology of the Cell*

Gustafson and Fromme

**Supplemental Table 1.** Yeast strains used in this study

<b>Name</b>	<b>Description</b>	<b>Source</b>
<b>SEY6210</b>	<i>MAT<math>\alpha</math> suc2-<math>\Delta</math>9 ura3-52 his3-<math>\Delta</math>200 leu2-3,112 lys2-801 trp1-<math>\Delta</math>901</i>	Robinson 1988 <sup>1</sup>
<b>SEY6210.1</b>	<i>MAT<math>\alpha</math> suc2-<math>\Delta</math>9 ura3-52 his3-<math>\Delta</math>200 leu2-3,112 lys2-801 trp1-<math>\Delta</math>901</i>	Robinson 1988 <sup>1</sup>
<b>BY4741<math>\alpha</math></b>	<i>MAT<math>\alpha</math> ura3-<math>\Delta</math>0 his3-<math>\Delta</math>1 leu2-<math>\Delta</math>0 lys2-<math>\Delta</math>0</i>	Brachmann 1998 <sup>2</sup>
<b>CFY578</b>	SEY6210 <i>Sec7-Mars::TRP1</i>	Richardson 2016 <sup>3</sup>
<b>CFY2376</b>	SEY6210 <i>Sec7-Mars::TRP1 Gea1-mNeonGreen::HIS3</i>	This study
<b>CFY2378</b>	SEY6210 <i>Sec7-Mars::TRP1 Gea2-mNeonGreen::HIS3</i>	This study
<b>CFY2490</b>	SEY6210.1 <i>Gea1-3xMars::TRP1 GFP-Vrg4</i>	This study
<b>CFY2503</b>	SEY6210.1 <i>Gea2-3xMars::TRP1 GFP-Vrg4</i>	This study
<b>CFY2872</b>	BY4741 $\alpha$ <i>gea1<math>\Delta</math>::KanMX gea2<math>\Delta</math>::HIS3 +pCF1248</i>	This study
<b>CFY2873</b>	BY4741 $\alpha$ <i>gea1<math>\Delta</math>::NatMX gea2<math>\Delta</math>::HIS3 arf1<math>\Delta</math>::KanMX +pCF1248</i>	This study

Name	Description	Vector	Source
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**Supplemental Table 2.** Plasmids used in this study

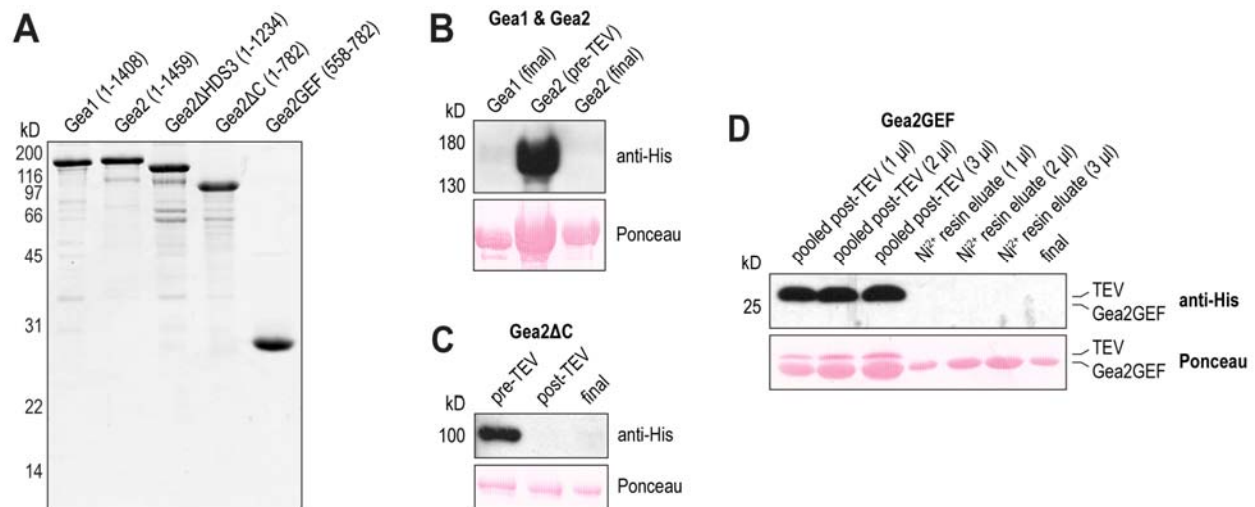
<b>pBCR314</b>	Sec7(203-2009) with a cleavable His <sub>6</sub> tag	pFastBacHT	Richardson 2012 <sup>4</sup>
<b>pMG005</b>	Gea2 FL (1-1459) with a cleavable His <sub>6</sub> tag	pET28	Richardson 2016 <sup>3</sup>
<b>pMG020</b>	Gea2ΔHDS3 (1-1196) with a cleavable His <sub>6</sub> tag	pET28	Richardson 2016 <sup>3</sup>
<b>pCF1299</b>	Gea2ΔC (1-782) with a cleavable His <sub>6</sub> tag	pET28	Richardson 2016 <sup>3</sup>
<b>pMG036</b>	Gea2GEF (558-782) with a cleavable His <sub>6</sub> tag	pET28	This study
<b>pCF1163</b>	Gea1 FL (1-1408) with a cleavable His <sub>6</sub> tag	pET28	Richardson 2012 <sup>4</sup>
<b>pArf1</b>	Arf1	pET3	Weiss 1989 <sup>5</sup>
<b>pCF1053</b>	Arf1ΔN17	pET28	Richardson 2012 <sup>4</sup>
<b>pCF1184</b>	Ar1	pET23	McDonold 2014 <sup>6</sup>
<b>pNMT1</b>	Nmt1	pCYC	Duronio 1990 <sup>7</sup>
<b>Ypt1-His<sub>7</sub></b>	Ypt1 with C-terminal His <sub>7</sub> tag and cleavable N-terminal GST tag	pGEX-6P	Gift from T. Bretscher
<b>pCM14</b>	Ypt6 with C-terminal His <sub>7</sub> tag and cleavable N-terminal GST tag	pGEX-6P	McDonold 2014 <sup>6</sup>
<b>pLT50</b>	Full-length Ypt1 with cleavable N-terminal GST tag	pGEX-6P	Thomas 2016 <sup>8</sup>
<b>pLT40</b>	Gdi1 with cleavable N-terminal GST tag	pGEX-6P	Thomas 2016 <sup>8</sup>
<b>pLT35</b>	Mrs6 with cleavable N-terminal His <sub>6</sub> tag	pET28	Thomas 2016 <sup>8</sup>
<b>pLT41</b>	Bet2 with cleavable N-terminal His <sub>6</sub> tag and Bet4	pCDF-Duet-1	Thomas 2016 <sup>8</sup>
<b>pCM10</b>	Gea1 FL (1-1408)-GFP driven by P <sub>G<sub>EA1</sub></sub>	pRS415	This study
<b>pCM09</b>	Gea1ΔHDS3 (1-1225)-GFP driven by P <sub>G<sub>EA1</sub></sub>	pRS415	This study
<b>pCF1312</b>	Gea1ΔC (1-761)-GFP driven by P <sub>G<sub>EA1</sub></sub>	pRS415	This study
<b>pMG001</b>	Gea2 FL (1-1459)-GFP driven by P <sub>G<sub>EA2</sub></sub>	pRS415	This study
<b>pMG002</b>	Gea2ΔHDS3 (1-1196)-GFP driven by P <sub>G<sub>EA2</sub></sub>	pRS415	This study
<b>pCF1313</b>	Gea2ΔC (1-766)-GFP driven by P <sub>G<sub>EA2</sub></sub>	pRS415	This study
<b>pCF1301</b>	Gea1-GFP driven by P <sub>G<sub>EA2</sub></sub>	pRS415	This study
<b>pCF1302</b>	Gea2-GFP driven by P <sub>G<sub>EA1</sub></sub>	pRS415	This study
<b>pCF1248</b>	Gea2 driven by P <sub>G<sub>EA2</sub></sub> (URA3 maintenance plasmid for shuffling strains)	pRS416	This study
<b>pRC2100</b>	GFP-Ypt1 driven by P <sub>YOP1</sub>	pRS415	Gift from R. Collins
<b>Ylplac211-iGFP-VRG4</b>	Integrating plasmid for GFP-VRG4 by two-step gene replacement	Ylplac211	Gift from B. Glick

**Supplemental Table 3.** Composition of liposomes used in this study

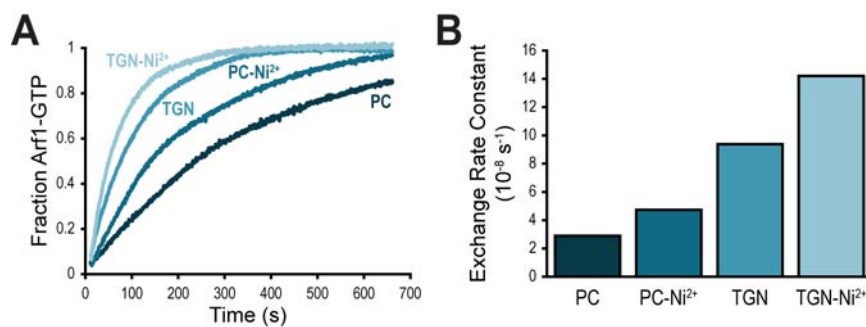
Lipid	PC	PC-Ni <sup>2+</sup>	TGN	TGN-Ni <sup>2+</sup>
	<i>Amount (Molar %)</i>			
DOPC	99	94	24	24
POPC			6	6
DOPE			7	7
POPE			3	3
DOPS			1	1
POPS			2	2
DOPA			1	1
POPA			2	2
PI			29	24
PI(4)P			1	1
CDP-DAG			2	2
PO-DAG			4	4
DO-DAG			2	2
Ceramide (C18)			5	5
Nickel-DOGS		5		5
Cholesterol			10	10
DiR	1	1	1	1

## Supplemental References

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4. Richardson, B. C., McDonold, C. M. & Fromme, J. C. The Sec7 Arf-GEF Is Recruited to the trans-Golgi Network by Positive Feedback. *Dev. Cell***22**, 799–810 (2012).
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6. McDonold, C. M. & Fromme, J. C. Four GTPases Differentially Regulate the Sec7 Arf-GEF to Direct Traffic at the trans-Golgi Network. *Dev. Cell***30**, 759–767 (2014).
7. Duronio, R. J. *et al.* Protein N-myristoylation in *Escherichia coli*: Reconstitution of a eukaryotic protein modification in bacteria. *Proc. Natl. Acad. Sci. U. S. A.***87**, 1506–1510 (1990).
8. Thomas, L. L. & Fromme, J. C. GTPase cross talk regulates TRAPP II activation of Rab11 homologues during vesicle biogenesis. *J. Cell Biol.***215**, 499–513 (2016).



**Supplemental Figure 1. GEF constructs used for biochemistry are pure and His<sub>6</sub> tags were successfully cleaved.** (A) Equal masses of full length Gea1 and Gea2, Gea2ΔHDS3, Gea2ΔC, and Gea2GEF were separated on an 8% polyacrylamide gel and visualized by Coomassie staining. Western blots with anti-His antibody against (B) Gea2 before TEV cleavage of the His<sub>6</sub> tag and final reagent samples of Gea1 and Gea2, (C) Gea2ΔC before and after TEV cleavage of the His<sub>6</sub> tag, as well as the final Gea2ΔC reagent, and (D) dilutions of fractions of Gea2GEF pooled after TEV cleavage, after incubation with and elution from Ni-NTA resin, and the final Gea2GEF reagent. For all western blots, Ponceau stains reveal protein on the membrane. Note that TEV protease is His<sub>6</sub>-tagged.



**Supplemental Figure 2. EDTA-induced Arf1 nucleotide exchange favors TGN over PC liposomes *in vitro*.** (A) Normalized traces showing EDTA exchange of Arf1 on PC, PC-Ni<sup>2+</sup>, TGN, and TGN-Ni<sup>2+</sup> liposomes. (B) Rates of Arf1 exchange determined from traces in (A).