## Supplementary file 4

**Table S7 Conformational states and transition kinetics from smGIET**

|  |  |  |
| --- | --- | --- |
| Protein | HOPS Vps33-yEGFP | HOPS Vps11-yEGFP |
| State  | L | M | H | L | M | H |
| *IG/I*0a | 0.25 ± 0.06 | 0.41 ± 0.08 | 0.69 ± 0.13 | 0.22 ± 0.05 | 0.36 ± 0.06 | 0.51 ± 0.08 |
| *h* (nm) b | 4.7(3.9 – 5.5) | 6.8(5.7 - 7.9) | 11.2(8.9 – 14.9) | 4.3(3.6 – 5.0) | 6.1(5.3 - 6.9) | 8.1(7.0 – 9.3) |
| *Occup.*(%)c | 30.9 ± 0.1 | 35.7 ± 0.2 | 33.5 ± 0.1 | 40.4 ± 0.2 | 38.0 ± 0.2 | 21.6 ± 0.1 |
| Transition counts and rate ***k*** (s-1)d | ***kL-M***: (471)1.05 ± 0.05***kL-H***: (65)0.15 ± 0.01 | ***kM-L***: (470)0.48 ± 0.02***kM-H***: (527)0.54 ± 0.02 | ***kH-L***: (77)0.13 ± 0.01***kH-M***: (534)0.87 ± 0.04 | ***kL-M***: (398)0.76 ± 0.04***kL-H***: (28)0.05 ± 0.01 | ***kM-L***: (392)0.53 ± 0.02***kM-H***: (267)0.36 ± 0.02 | ***kH-L***: (34)0.10 ± 0.01***kH-M***: (262)0.73 ± 0.04 |
| *ΔG (kBT)e* | +0.78 | 0 | +0.48 | +0.36 | 0 | +0.71 |

*a*: mean ± s.d. based on *IG* of the Gaussian fits in single molecule intensity distribution on graphene. *I*0 is the mean value of Gaussian fit on glass (Fig S15).*b*: *h* is the height of NB-labeled HOPS on the membrane. Values in brackets are the boundary of *h*. *c*: mean ± s.e.m. of state occupancy based on observations of N = 101615 for Vps33, N = 84218 for Vps11, respectively. *d*: Transition counts are shown in brackets. Transition rates are presented as mean ± s.e.m. *e*: Free energy between the states calculated by $ΔG=-k\_{B}TlnK$, in which $k\_{B}$ is the Boltzmann constant. *T* is the absolute temperature. *K* is the equilibrium constant calculated as the ratio of forward and backward transition rates between M and the target state.