SUPPLEMENTARY ONLINE DATA Polyphenols differentially inhibit degranulation of distinct subsets of vesicles in mast cells by specific interaction with granule-type-dependent SNARE complexes

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(a) All 32 phenolic compounds used representing 11 subgroups and their abbreviations. (b and c) Inhibitory effect of polyphenols on β -hexosaminidase and histamine release from the RBL-2H3 cells. RBL-2H3 cells were treated with or without 10 μ M polyphenols. Results are means \pm S.D. from five independent experiments. Dense area of (b) is shown enlarged in (c) for clarity.

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Figure S2 Effect of polyphenols on mast cell degranulation is independent of stimulation method

 β -Hexosaminidase release from RBL-2H3 cells was stimulated by either Fc_ERI cross-linking or calcium ionophore A23187 in the presence or absence of polyphenols.



Figure S3 The lipid mixing ability of various SN25-containing SNARE complexes

SNAP-25 or Syn1a could induce lipid mixing with various VAMPs, but these proteins are not expressed in mast cells. SN25, 25 kDa synaptosome-associated protein; Vp, vesicle-associated membrane protein.



Figure S4 Inhibitory effect of 32 polyphenols on membrane fusion driven by three SNARE complexes

(a) SNAP-23/Syn4/VAMP2 compared with SNAP-23/Syn4/VAMP8. (b) SNAP-23/Syn4/VAMP2 compared with SNAP-25/Syn1/VAMP2. Dense areas of (a and b) are shown enlarged in (c) and (d) respectively for better resolution. SN23/25, 23/25 kDa synaptosome-associated protein; Vp, vesicle-associated membrane protein.





Soluble fragments of SNARE proteins (Vp2S and Vp8S) and t-SNARE complex were mixed with polyphenols and UV spectra were measured. Abs., absorbance; CY, cyanidin; DL, delphinidin; SN23/25, 23/25 kDa synaptosome-associated protein; Vp, vesicle-associated membrane protein.



Figure S6 Dense areas of Figure 4 of the main text are shown enlarged



Figure S7 Determination of IC₅₀ values of polyphenols for histamine and β-hexosaminidase release from RBL-2H3 cells

Table S1	Primers and condition for PCR analysis of RBL-2H3 cells
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F, forward; R, reverse.

Туре	Gene	Primer type	Primer sequence $(5' \rightarrow 3')$	Annealing temperature (°C)	Number of cycles	Amplicon size (bp)
v-SNARES	VAMP2	F	ATGTCGGCTACCGCTGCCAC	58	28	348
		R	ATCGTTTACTTCAGCACT			
	VAMP8	F	ATGGAGGCCAGTGGGAGTGC	58	28	300
		R	GGCACCATCCCCACT			
	VAMP8S	F	ATGGAGGCCAGTGGGAGTGC	58	28	222
		R	GTTCTGGTGGAAGAATGTGAAG			
	VAMP4	F	ATGCCTCCCAAGTTCAAGCG	58	30	423
		R	TAGTTGTGAAATACCGTACT			
	VAMP7	F	ATGGCCATTCTTTTTGCCGTTG	58	28	660
		R	GGCCAAGCTGTGTGAAGAAA			
t-SNAREs	SNAP-23	F	ATGGATGATCTATCACCAGA	52	30	630
		R	AGAGCAAAGAAACTCATTGACAGC			
	SNAP-25	F	ATGGCCGAGGACGCAGACAT	52	40	618
		R	AAGATGCTGGGAAGTGGT			
	Syntaxin 1a	F	ATGAAGGACCGAACCCAGGA	56	40	864
		R	ACCATCGGGGGCATCTTTGGA			
	Syntaxin 4	F	ATGCGCGACAGGACCCATGA	56	40	894
		R	CATCACCATAACCGTTGGA			
	Syntaxin 5	F	ATGTCCTGCCGGGATCGGAC	56	40	903
		R	ATCTTTGTGGTCTTCCTTGCC			
	β -Actin	F	ATGGGTCAGAAGGACTCCTACG	58	18	470
		R	CATTGTGATGGACTCCGGAGA			

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