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	Allergen - Scientific Name	Swiss-Prot ID (AccNum)	PDB ID	HSSP AccNum (ID)
1	Aca s 13	FABP_ACASI (O76821)		P04117 (1LID)
2	Act c 1	ACTN_ACTCH (P00785)	2ACT, 1AEC, 1F3Q, 1FOG, 1FTV	
3	Aln g 1	MPAG_ALNGL (P38948)		P15494 (1BV1)
4	Aln g 4	POC4_ALNGL (O81701)		P02593 (1AK8)
5	Alt a 10	DHAL_ALTAL (P42041)		P05091 (1CW3)
6	Alt a 11	ENO_ALTAL (Q9HDT3)		P00924 (1ONE)
7	Amb a 3	MPA3_AMBEL (P00304)		P00303 (2CBP)
8	Amb a 5	MPA5_AMBEL (P02878)		P10414 (1BBG)
9	Amb a 6	NLT6_AMBAR (O04004)		P19656 (1MZM)
10	Amb p 5	MP5A_AMBPS (P43174), MP5B_AMBPS (P43175)		P10414 (1BBG), P10414 (1BBG)
11	Amb t 5	MPT5_AMBTR (P10414)	1BBG, 2BBG, 3BBG	
12	Api g 1	ALL1_APIGR (P49372)		P15494 (1BV1)
13	Api g 2	ALL2_APIGR (P92918)		O24248 (1E09)
14	Api g 3	CB23_APIGR (P92919)		P04083 (1BO9)
15	Api g 4	PROF_APIGR (Q9XF37)		P25816 (1CQA)
16	Api m 1	PA2_APIME (P00630)	1POC	
17	Api m 2	HUGA_APIME (Q08169)	1FCQ, 1FCU, 1FCV	
18	Api m 3	MEL_APIME (P01501)	2MLT, 1BH1	
19	Ara h 1	AL11_ARAHY (P43237), AL12_ARAHY (P43238)		P50477 (1CAU), P50477 (1CAU)
20	Ara h 5	PROF_ARAHY (Q9SQI9)		P25816 (1CQA)
21	Ara t 8	PRO1_ARATH (Q42449)	3NUL, 1A0K, 1PLM	
22	Asp f 1	RNMG_ ASPRE (P04389)	1AQZ, 1JBR, 1JBS, 1JBT, 1JPU, 1JQ5,	
23	Asp f 3	PM20_ASPFU (O43099)		P30044 (1H4O)
24	Asp f 6	SODM_ASPFU (Q92450)	1KKC	
25	Ber e 1	2SS_BEREX (P04403)	1GYS	
26	Bet v 1	BV1A_BETVE (P15494), BV1B_BETVE (P45431), BV1C_BETVE (P43176), BV1D_BETVE (P43177), BV1E_BETVE (P43178), BV1F_BETVE (P43179), BV1G_BETVE (P43180), BV1J_BETVE (P43183), BV1K_BETVE (P43184), BV1L_BETVE (P43185), BV1M_BETVE (P43186)	1BTV, 1BV1, 1B6F, 1FSK, 1QMR, 1FM4	P15494 (1BV1)
27	Bet v 2	PROF_BETVE (P25816)	1CQA	
28	Bet v 3	ALL3_BETVE (P43187)		P02593 (1CDM)
29	Bet v 4	POC4_BETVE (Q39419)		P02593 (1CDM)
30	Bet v 7	CYPH_BETVE (P81531)		P05092 (2CPL)
31	Bla g 5	GTS1_BLAGE (O18598)		P24472 (1GUK)
32	Bla g II	ASP2_BLAGE (P54958)		P00797 (2REN)
33	Blo t 12	BT12_BLOTA (Q17282)		P10724 (1BD0)
34	Blo t 13	FABP_BLOTA (Q17284)		P04117 (1LID)
35	Bom t 1	PA2_BOMTE (P82971)		P00630 (1POC)
36	Bos d 2	ALL2_BOVIN (Q28133)	1BJ7	
37	Bos d 3	S107_BOVIN (Q28050)		P31151 (1PSR)
38	Bos d 4	LCA_BOVIN (P00711)	1HFZ, 1F6R, 1F6S	
39	Bos d 5	LACB_BOVIN (P02754)	1BEB, 1BSO, 1BSQ, 1BSY, 2BLG, 3BLG, 1B00, 1CJ5, 1B8E, 1DV9, 1GX8, 1GX9, 1GXA, 1MFH, 1QG5	
40	Bos d 6	ALBU_BOVIN (P02769)		P02768 (1E7B)

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41	Bra n 1	POC1_BRARA (Q42470)		P02593 (1CDL)
42	Bra n 2	POC2_BRARA (Q39406)		P02588 (1AVS)
43	Bra r 1	POC1_BRARA (Q42470)		P02593 (1CDL)
44	Bra r 2	POC2_BRARA (Q39406)		P02588 (1AVS)
45	Bra r 2	CHAL_BRARA (P81729)		P28814 (1BW4)
46	Can f 3	ALBU_CANFA (P49822)		P02768 (1E7B)
47	Cand b 2	PMPA_CANBO (P14292), PMPB_CANBO (P14293)		P30044 (1HD2), P30044 (1HD2)
48	Car b 1	MPA1_CARBE (P38949), MPA2_CARBE (P38950)		P15494 (1BV1), P15494 (1BV1)
49	Cav p 2	AVP2_CAVPO (P83508)	1I49, 1I4D, 1I4L, 1I4T	
50	Cla h 3	DHAL_CLAHE (P40108)		P05091 (1CW3)
51	Cla h 4	HS70_CLAHE (P40918)		P19120 (3HSC)
52	Cla h 6	ENO_CLAHE (P42040)		P00924 (4ENL)
53	Cop c 2	THIO_COPCM (Q9UW02)		P80028 (1TOF)
54	Cor a 1	MPAA_CORAV (Q08407)		P15494 (1BV1)
55	Cry j 2	MPA2_CRYJA (P43212)		P26509 (1BHE)
56	Cyn d 1	MPC1_CYNDA (O04701)		P43214 (1WHO)
57	Cyn d 7	POC7_CYNDA (P94092)		P25070 (1AVJ)
58	Cyn d 12	PROF_CYNDA (O04725)		P25816 (1CQA)
59	Dac g 2	MPG2_DACGL (Q41183)		P43214 (1WHO)
60	Dac g 3	MPG3_DACGL (P93124)		P43214 (1WHO)
61	Dac g 4	MPG4_DACGL (P82946)	1DXY	
62	Dau c 1	DAU1_DAUCA (O04298)	1BGL, 1BGM, 1DP0, 1F49, 1F4A, 1F4H, 1GHO, 1HN1, 1JYN, 1JYV, 1JYW, 1JYX, 1JYY, 1JYZ, 1JZ0, 1JZ1, 1JZ2, 1JZ3, 1JZ4, 1JZ5, 1JZ6, 1JZ7, 1JZ8	O24248 (1E09)
63	Der f 1	MMAL_DERFA (P16311)		P07688 (1QDQ)
64	Der f 2	ALL2_DERFA (Q00855)	1AHK, 1AHM	
65	Der f 3	DEF3_DERFA (P49275)		P00763 (1DPO)
66	Der f 6	DEF6_DERFA (P49276)		P00763 (1DPO)
67	Der p 1	MMAL_DERPT (P08176)		P07858 (1HUC)
68	Der p 2	ALL2_DERPT (P49278)	1A9V, 1KTJ	
69	Der p 3	DER3_DERPT (P39675)		P00763 (1DPO)
70	Der p 8	GTM1_DERPT (P46419)		P09488 (1GTU)
71	Dol a 5	VA5_DOLAR (Q05108)		P04284 (1CFE)
72	Dol m 1	PA11_DOLMA (Q06478), PA12_DOLMA (P53357)		P54318 (1BU8), P54318 (1BU8)
73	Dol m 5	VA52_DOLMA (P10736), VA53_DOLMA (P10737)		P04284 (1CFE)
74	Equ c 1	ALL1_HORSE (Q95182)	1EW3	
75	Equ c 3	ALBU_HORSE (P35747)		P02768 (1E7B)
76	Eur m 1	EUM1_EURMA (P25780)		P07688 (1QDQ)
77	Eur m 2	ALL2_EURMA (Q9TZZ2)		P49278 (1A9V)
78	Eur m 3	EUM3_EURMA (O97370)		P00763 (1DPO)
79	Fel d 2	ALBU_FELCA (P49064)		P02768 (1E7B)
80	Gad c 1	PRVB_GADCA (P02622)		P02618 (1CDP)
81	Gad m 1	PRVB_GADMO (Q90YK9)		P02618 (1B8C)
82	Gal d 1	IOVO_CHICK (P01005)		P05586 (4OVO)

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83	Gal d 2	OVAL_CHICK (P01012)	1OVA, 1JTI, 1VAC	
84	Gal d 3	TRFE_CHICK (P02789)	1NNT, 1OVT, 1AIV, 1IEJ, 1JL4, 1NFT,	
85	Gal d 4	LYC_CHICK (P00698)	2LYM, 3LYM, 4LYM, 5LYM, 1LYS, 3LYT, 4LYT, 5LYT, 6LYT, 1LYZ, 2LYZ, 3LYZ, 4LYZ, 5LYZ, 6LYZ, 7LYZ, 8LYZ, 1LZH, 2LZH, 1LZT, 2LZT, 3LZT, 4LZT, 3HFL, 3HFM, 1HEL, 1HEM, 1HEN, 1HEO, 1HEP, 1HEQ, 1HER, 1HEW, 1LMA, 1RCM, 1LSA, 1LSB, 1LSC, 1LSD, 1LSE, 1LSF, 1LSM, 1LS	P04036 (1DRW)
86	Gal d 5	ALBU_CHICK (P19121)		P02768 (1E7B)
87	Gly m 1	HPSE_SOYBN (P24337)	1HYP	
88	Gly m 3	PRO1_SOYBN (O65809), PRO2_SOYBN (O65810)		P25816 (1CQA), P25816 (1CQA)
89	Hel a 2	PROF_HELAN (O81982)		P25816 (1CQA)
90	Hev b 6	HEVE_HEVBR (P02877)	1HEV	
91	Hev b 8	PRO1_HEVBR (O65812), PRO2_HEVBR (Q9STB6), PRO3_HEVBR (Q9M7N0), PRO4_HEVBR (Q9M7M9), PRO5_HEVBR (Q9M7M8), PRO6_HEVBR (Q9LEI8)	1G5U	P25816 (1CQA), P25816 (1CQA), P25816 (1CQA), P25816 (1CQA), P25816 (1CQA)
92	Hev b 9	ENO1_HEVBR (Q9LEJ0), ENO2_HEVBR (Q9LEI9)		P56252 (1PDZ), P56252 (1PDZ)
93	Hol l 1	MPH1_HOLLA (P43216)		P43214 (1WHO)
94	Hor v 1	IAA1_HORVU (P16968)		P01085 (1HSS)
95	Jun a 3	PRR3_JUNAS (P81295)		P33679 (1DU5)
96	Jun o 2	POC2_JUNOX (O64943)		P02593 (1CDM)
97	Lep d 1	ALL1_LEPDS (P80384)		Q00855 (1AHK)
98	Lep d 13	FABP_LEPDS (Q9U5P1)		P04117 (1LID)
99	Lol p 1	MPL1_LOLPR (P14946)		P43214 (1WHO)
100	Lol p 2	MPL2_LOLPR (P14947)		P43214 (1WHO)
101	Lol p 3	MPL3_LOLPR (P14948)		P43214 (1WHO)
102	Mal d 1	MAL1_MALDO (P43211)		Q24248 (1E09)
103	Mal d 2	TP1A_MALDO (Q9FSG7)		P33679 (1DU5)
104	Mal d 3	NLTP_MALDO (Q9M5X7)		P19656 (1MZM)
105	Mal d 4	PRO1_MALDO (Q9XF40), PRO2_MALDO (Q9XF41), PRO3_MALDC		P25816 (1CQA), P25816 (1CQA), P25816 (1CQA)
106	Mal f 3	MLF3_MALFU (P56578)		P30044 (1HD2)
107	Mer a 1	PROF_MERAN (O49894)		P25816 (1CQA)
108	Mus m 1	MUP6_MOUSE (P02762)	1MUP, 1I04, 1I05, 1I06	
109	Ole e 2	PRO1_OLEEU (O24169), PRO2_OLEEU (O24170), PRO3_OLEEU (O24171)		P25816 (1CQA), P25816 (1CQA), P25816 (1CQA)
110	Ole e 3	POC3_OLEEU (O81092)		P02588 (1AVS)
111	Ole e 8	ALL8_OLEEU (Q9M7R0)		P02593 (1CTR)
112	Ory s 1	MPO1_ORYSA (Q40638)		P43214 (1WHO)
113	Par j 1	NL11_PARJU (P43217), NL12_PARJU (O04404), NL13_PARJU (Q40905)		P19656 (1MZM), P19656 (1MZM), P19656
114	Par j 2	NL21_PARJU (P55958), NL22_PARJU (O04403)		P23096 (1RZL), P23096 (1RZL)
115	Par j 3	PRO1_PARJU (Q9XG85), PRO2_PARJU (Q9T0M8)		P25816 (1CQA), P25816 (1CQA)
116	Pen c 19	HS70_PENCI (Q92260)		P19120 (3HSC)
117	Per a 3	CRPI_PERAM (Q25641)		P04253 (1OXY)
118	Pha a 1	MPA1_PHAAQ (Q41260)		P43214 (1WHO)

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119	Phl p 1	MPP1_PHLPR (P43213)	1N10	
120	Phl p 2	MPP2_PHLPR (P43214)	1WHO, 1WHP, 1BMW	
121	Phl p 5b	MP5B_PHLPR (Q40963)	1L3P	
122	Phl p 6	MPP6_PHLPR (P43215)	1NLX	
123	Phl p 7	POC7_PHLPR (O82040)		P25070 (1AVJ)
124	Phl p 11	PRO1_PHLPR (P35079), PRO2_PHLPR (O24650), PRO3_PHLPR (O24282)		P25816 (1CQA), P25816 (1CQA), P25816 (1CQA)
125	Pol a 1	PA1_POLAN (Q9U6W0)		P00591 (1ETH)
126	Pol a 5	VA5_POLAN (Q05109)		P04284 (1CFE)
127	Pol d 5	VA5_POLDO (P81656)		P04284 (1CFE)
128	Pol e 5	VA5_POLEX (P35759)		P04284 (1CFE)
129	Pol f 5	VA5_POLFU (P35780)		P04284 (1CFE)
130	Pru ar 1	PRU1_PRUAR (O50001)		O24248 (1E09)
131	Pru ar 3	NLT1_PRUAR (P81651)		P19656 (1MZM)
132	Pru av 1	PRU1_PRUAV (O24248)	1E09	
133	Pru av 3	NLTP_PRUAV (Q9M5X8)		P19656 (1MZM)
134	Pru av 4	PROF_PRUAV (Q9XF39)		P25816 (1CQA)
135	Pru d 3	NLT1_PRUDO (P82534)		P19656 (1MZM)
136	Pru p 3	NLT1_PRUPE (P81402)		P19656 (1MZM)
137	Pyr c 1	PYR1_PYRCO (O65200)		O24248 (1E09)
138	Pyr c 3	NLTP_PYRCO (Q9M5X6)		P19656 (1MZM)
139	Pyr c 4	PROF_PYRCO (Q9XF38)		P25816 (1CQA)
140	Rat n 1	MUP_RAT (P02761)	2A2G, 2A2U	
141	Sal s 1	PRV1_SALSA (Q91482), PRV2_SALSA (Q91483)		P02619 (2PVB), P02619 (2PVB)
142	Sola t 4	SPI7_SOLTU (P30941)	1AOI, 1M18, 1M19	
143	Sol i 3	VA3_SOLIN (P35778)		P04284 (1CFE)
144	Sol r 3	VA3_SOLRI (P35779)		P04284 (1CFE)
145	Syr v 3	POC3_SYRVU (P58171)		P02588 (1AVS)
146	The c 1	PRVB_THECH (Q90YK7)		P02618 (1B8C)
147	Tyr p 2	ALL2_TYRPU (O02380)		Q00855 (1AHK)
148	Ves f 5	VA5_VESFL (P35783)		P04284 (1CFE)
149	Ves g 5	VA5_VESGE (P35784)		P04284 (1CFE)
150	Ves m 1	PA1_VESMC (P51528)		P54318 (1BU8)
151	Ves m 5	VA5_VESMC (P35760)		P04284 (1CFE)
152	Ves p 5	VA5_VESPE (P35785)		P04284 (1CFE)
153	Ves s 5	VA5_VESSQ (P35786)		P04284 (1CFE)
154	Ves v 1	PA1_VESVU (P49369)		P29183 (1HPL)
155	Ves v 5	VA5_VESVU (Q05110)	1QNX	
156	Ves vi 5	VA5_VESVI (P35787)		P04284 (1CFE)
157	Vesp c 5	VA51_VESCR (P35781), VA52_VESCR (P35782)		P04284 (1CFE), P04284 (1CFE)
158	Zea m 1	EXBA_MAIZE (P58738), EXBB_MAIZE (Q07154)		P43214 (1WHO)
159	Zea m 14	NLTP_MAIZE (P19656)	1MZL, 1MZM, 1AFH, 1FK0, 1FK1, 1FK2, 1FK3, 1FK4, 1FK5, 1FK6, 1FK7	