

Characterising Institute code ITADiSTA

Country of characterisation Italy

Farm or Institute address

Latitude

Longitude

Elevation

Evaluators name Prof. Filippo d'Antuono

Evaluation environment Field

Institute code	Accession number	Sowing date	Anthocyanin in petiole	Petiole hairiness	Leaf hairiness	Leaf dissection
FRAINH	1-INH	2001	1.0	1.3	1.4	6.4
FRAINH	2-INH	2001	1.0	1.3	1.4	6.3
FRAINH	3-INH	2001	1.0	1.3	1.4	6.9
FRAINH	4-INH	2001	1.1	1.2	1.2	6.4
FRAINH	5-INH	2001	1.1	1.3	1.6	6.6
FRAINH	6-INH	2001	3.7	1.5	1.9	5.7
FRAINH	7-INH	2001	1.0	1.3	1.5	6.6
FRAINH	8-INH	2001	1.0	1.3	1.4	6.4
FRAINH	9-INH	2001	1.0	1.3	1.2	6.7
FRAINH	10-INH	2001	1.0	1.1	1.1	6.9
Standard	Parmex	2001	1.0	1.1	1.4	6.1
Standard	Rubrovitamina	2001	1.0	1.5	1.5	6.3
DEU001	BAZ-59153	2001	1.0	1.4	1.6	6.9
DEU001	BAZ-69565	2001	1.0	1.1	1.2	6.6
DEU001	BAZ-69563	2001	1.0	1.3	1.5	6.3
DEU001	BAZ-56374	2001	1.0	1.3	1.3	5.8
DEU001	BAZ-56367	2001	1.0	1.6	1.7	6.4
DEU001	BAZ-56368	2001	1.0	1.6	1.8	6.4
DEU001	BAZ-56376	2001	1.0	1.3	1.6	6.6
DEU001	BAZ-56375	2001	1.0	1.5	1.5	6.4
DEU001	BAZ-55504	2001	1.0	1.3	1.3	6.5
DEU001	BAZ-46588	2001	1.0	1.8	1.9	6.4
DEU001	BAZ-65695	2001	1.0	1.4	1.6	6.6
DEU001	BAZ-62634	2001	1.0	1.1	1.2	6.7
DEU001	BAZ-55505	2001	1.0	1.1	1.3	6.2
DEU001	BAZ-46591	2001	1.0	1.4	1.6	6.8
DEU001	BAZ-59152	2001	1.0	1.3	1.6	6.3
DEU001	BAZ-62633	2001	1.0	1.5	1.6	6.2
DEU001	BAZ-46587	2001	1.0	1.3	1.8	6.5
DEU001	BAZ-46584	2001	1.0	1.2	1.1	6.9
DEU001	BAZ-69566	2001	1.0	1.2	1.8	6.3
DEU001	BAZ-52136	2001	1.0	2.2	1.8	5.4
Standard	Amsterdam	2001	1.0	1.0	1.2	6.5
Standard	Bolero	2001	1.0	1.2	1.2	6.8
GBRHRIGRU	HRI-3931	2001	1.0	2.3	1.9	5.2
GBRHRIGRU	HRI-3933	2001	1.0	1.3	1.7	6.1
GBRHRIGRU	HRI-3937	2001	1.0	1.4	1.5	6.8
GBRHRIGRU	HRI-4003	2001	1.0	1.5	1.4	6.2
GBRHRIGRU	HRI-4007	2001	1.0	1.5	1.9	6.6
GBRHRIGRU	HRI-4011	2001	1.0	2.8	4.6	4.5
GBRHRIGRU	HRI-4012	2001	1.1	3.3	4.0	4.5
GBRHRIGRU	HRI-4013	2001	1.0	1.9	2.0	6.2

GBRHRIGRU	HRI-5779	2001	1.0	1.3	1.4	6.3
GBRHRIGRU	HRI-6183	2001	1.0	1.3	1.2	6.8
GBRHRIGRU	HRI-6184	2001	1.0	1.3	1.5	5.7
GBRHRIGRU	HRI-6486	2001	1.0	1.2	1.1	7.0
GBRHRIGRU	HRI-6497	2001	1.0	1.8	2.0	6.3
GBRHRIGRU	HRI-6519	2001	1.0	1.0	1.2	7.2
GBRHRIGRU	HRI-6682	2001	1.0	2.1	2.1	5.7
GBRHRIGRU	HRI-6686	2001	2.2	1.2	1.5	6.5
GBRHRIGRU	HRI-6688	2001	1.0	1.2	1.4	6.3
GBRHRIGRU	HRI-6689	2001	1.3	1.8	1.5	5.1
GBRHRIGRU	HRI-6756	2001	1.0	1.6	1.1	5.9
GBRHRIGRU	HRI-6764	2001	1.0	1.3	1.4	6.7
GBRHRIGRU	HRI-7170	2001	1.0	5.4	7.0	3.1
GBRHRIGRU	HRI-7172	2001	1.0	1.2	1.5	6.2
GBRHRIGRU	HRI-7890	2001	1.0	1.1	1.1	7.0
GBRHRIGRU	HRI-7902	2001	1.0	1.7	2.0	6.5
GBRHRIGRU	HRI-7904	2001	1.6	1.5	1.6	6.9
GBRHRIGRU	HRI-8684	2001	1.0	1.3	1.2	6.1
GBRHRIGRU	HRI-8685	2001	1.0	1.3	1.2	6.0
GBRHRIGRU	HRI-8715	2001	1.0	2.4	2.7	3.8
GBRHRIGRU	HRI-8716	2001	1.0	2.2	2.1	4.1
GBRHRIGRU	HRI-8727	2001	1.0	1.0	1.0	6.0
GBRHRIGRU	HRI-9808	2001	1.0	1.6	1.7	6.7
GBRHRIGRU	HRI-10146	2001	-999	-999	-999	-999
GBRHRIGRU	HRI-10149	2001	1.1	1.6	1.6	6.3
GBRHRIGRU	HRI-10156	2001	1.0	1.7	1.3	5.7
GBRHRIGRU	HRI-10165	2001	1.8	2.5	1.6	5.3
GBRHRIGRU	HRI-10228	2001	2.3	1.1	1.1	4.4
GBRHRIGRU	HRI-10264	2001	1.0	2.1	2.0	6.5
GBRHRIGRU	HRI-10522	2001	1.0	1.5	1.3	6.2
GBRHRIGRU	HRI-10524	2001	2.4	2.1	5.9	3.3
GBRHRIGRU	HRI-11201	2001	3.8	2.3	2.0	5.4
GBRHRIGRU	HRI-11261	2001	1.0	1.5	1.3	6.1
Standard	Autumn king	2001	1.0	1.0	1.6	6.5
	News F1	2002	1.0	1.0	1.1	6.8
FRAINH	Russe n°7 - INH 11	2002	1.0	1.5	1.2	5.4
FRAINH	Amsterdamer- INH 12	2002	1.0	1.2	1.5	6.8
FRAINH	Jere - INH 13	2002	2.7	1.1	1.2	6.5
FRAINH	Parijse Markt 2 - INH 14	2002	1.0	1.5	1.6	6.7
FRAINH	Boev - INH 15	2002	1.0	1.2	1.3	6.9
FRAINH	Eysines -INH 16	2002	1.0	1.6	1.7	6.9
FRAINH	Rote Riesen - INH 17	2002	1.0	1.3	1.5	6.8
FRAINH	St Valery Sileban-INH 18	2002	1.0	1.6	1.9	6.9
FRAINH	Russe n°3 - INH 19	2002	1.0	1.1	1.3	6.8
FRAINH	Nantaiser 58 - INH 20	2002	1.0	1.1	1.3	6.9
Standard	Parmex	2002	1.0	1.0	1.1	6.8
	Loc ITA 1	2002	1.1	1.7	1.9	6.8
DEU001	56341-BAZ	2002	1.0	1.9	2.1	7.4
DEU001	56335-BAZ	2002	1.0	1.3	1.8	7.0
Standard	Amsterdam	2002	1.0	1.1	1.9	6.8
Standard	Nikki F1	2002	1.0	1.1	1.8	6.8
GBRHRIGRU	3838 - HRI	2002	1.0	1.0	1.5	7.4
GBRHRIGRU	3936 - HRI	2002	1.3	1.7	1.8	6.7
GBRHRIGRU	3966 - HRI	2002	1.0	1.1	1.1	6.8
GBRHRIGRU	3998 - HRI	2002	1.1	1.4	1.7	6.9
GBRHRIGRU	5784 - HRI	2002	1.0	1.1	1.5	7.0

GBRHIGRU	6070 - HRI	2002	1.0	1.0	1.1	6.5
GBRHIGRU	6102 - HRI	2002	1.0	1.1	1.4	6.7
GBRHIGRU	6755 - HRI	2002	1.1	1.9	1.7	6.6
GBRHIGRU	6760 - HRI	2002	2.1	1.2	1.3	6.7
GBRHIGRU	7174 - HRI	2002	1.0	2.1	2.1	6.8
GBRHIGRU	7301 - HRI	2002	5.3	1.1	1.1	6.4
GBRHIGRU	7801 - HRI	2002	1.0	1.5	1.6	7.1
GBRHIGRU	7893 - HRI	2002	1.0	1.0	1.3	6.8
GBRHIGRU	8080 - HRI	2002	1.0	1.0	1.2	6.7
GBRHIGRU	8081 - HRI	2002	1.0	1.1	1.7	6.8
GBRHIGRU	8095 - HRI	2002	1.0	1.2	1.9	6.8
GBRHIGRU	8116 - HRI	2002	1.0	1.3	1.7	6.9
GBRHIGRU	8125 - HRI	2002	1.0	1.1	1.4	6.4
GBRHIGRU	8394 - HRI	2002	1.1	1.2	1.6	6.8
GBRHIGRU	10168 - HRI	2002	1.5	2.5	2.5	6.1
GBRHIGRU	10176 - HRI	2002	2.8	1.7	1.5	7.0
GBRHIGRU	10194 - HRI	2002	1.4	3.5	3.3	6.1
GBRHIGRU	10220 - HRI	2002	1.6	3.3	3.4	5.5
GBRHIGRU	10225 - HRI	2002	3.4	2.1	1.9	6.2
GBRHIGRU	10233 - HRI	2002	1.1	1.2	1.6	7.1
GBRHIGRU	10237 - HRI	2002	2.2	2.1	2.4	6.8
GBRHIGRU	10246 - HRI	2002	1.1	1.4	1.6	6.7
GBRHIGRU	10261 - HRI	2002	1.0	1.0	1.2	6.7
GBRHIGRU	10305 - HRI	2002	1.1	1.3	1.8	7.0
GBRHIGRU	10385 - HRI	2002	1.1	3.3	3.0	7.1
GBRHIGRU	10468 - HRI	2002	1.3	4.0	3.8	5.5
GBRHIGRU	10520 - HRI	2002	1.0	1.6	1.5	7.1
GBRHIGRU	10521 - HRI	2002	4.1	2.9	2.9	7.0
GBRHIGRU	11163 - HRI	2002	1.0	1.3	1.7	6.8
GBRHIGRU	11169 - HRI	2002	1.0	1.1	1.2	7.1
GBRHIGRU	11503 - HRI	2002	1.0	1.0	1.1	6.8
GBRHIGRU	12869 - HRI	2002	1.2	2.0	1.5	7.7
Standard	Autumn King	2002	1.0	1.1	1.5	6.7
	Loc ITA 2	2002	1.3	1.4	1.6	7.2
REGNGB	2399 - NGB	2002	1.2	1.1	1.3	6.9
REGNGB	7748 - NGB	2002	1.0	1.2	1.8	6.9
REGNGB	13942 - NGB	2002	-999	-999	-999	-999
REGNGB	13943 - NGB	2002	-999	-999	-999	-999
REGNGB	13945 - NGB	2002	1.0	1.0	1.5	6.9
REGNGB	13946 - NGB	2002	1.0	1.2	1.7	6.7
REGNGB	13949 - NGB	2002	1.0	1.1	1.7	6.8
REGNGB	13955 - NGB	2002	1.0	1.6	2.1	6.6
REGNGB	13951 - NGB	2002	1.0	1.1	1.4	6.4
REGNGB	13636 - NGB	2002	1.0	1.2	1.5	6.9
GBRHIGRU	5798-HRI	2002	1.3	1.2	1.7	6.9

Foliage width (crown)	Root length	Root diameter	diameter relative to total	Root branching	Root shape in longitudinal section	Root shoulder shape	Root tapering	Root tip shape
5.2	5.6	4.7	6.5	1.4	3.3	2.8	0.7	2.5
4.1	6.2	4.9	6.0	0.6	3.2	2.5	1.0	2.8
4.4	3.0	3.7	5.7	1.5	2.6	2.6	1.2	2.8
5.0	6.0	4.8	6.3	1.1	3.2	2.8	0.9	2.6
4.5	4.6	4.0	5.5	1.1	3.6	2.7	0.5	2.2
4.9	4.3	5.6	6.5	1.4	3.3	3.1	1.2	2.4
4.6	4.5	4.5	5.6	1.2	3.1	2.6	1.0	2.8
5.5	5.1	5.5	6.0	1.3	3.0	2.4	1.6	2.7
5.4	6.1	4.7	7.4	1.4	3.0	2.4	1.3	2.9
4.6	4.8	4.2	4.7	0.5	3.7	2.6	0.8	2.2
4.7	2.6	5.7	5.0	1.2	1.7	2.4	0.4	1.9
5.1	4.9	3.9	6.4	1.3	3.2	2.8	1.1	2.8
4.6	4.7	4.0	5.3	0.6	3.1	2.6	1.2	2.9
4.4	4.7	3.5	4.9	0.9	3.5	2.4	0.8	2.4
4.5	3.9	4.3	6.5	1.1	3.0	2.5	1.9	2.7
4.9	4.6	4.9	6.2	1.3	2.4	2.4	1.8	2.7
4.6	4.9	4.8	5.6	1.4	3.3	2.8	1.1	2.5
5.7	4.4	6.2	6.4	1.5	2.4	2.4	2.3	2.6
4.6	5.4	4.4	6.0	1.4	3.1	2.5	1.1	2.6
4.6	3.7	5.6	6.6	1.3	2.0	2.4	1.4	1.8
4.6	4.1	3.7	6.2	0.7	3.3	2.6	0.9	2.7
4.7	3.9	4.3	5.4	1.3	3.0	2.4	1.5	3.0
4.8	4.3	4.7	6.2	1.0	3.1	2.9	1.3	2.8
4.7	4.3	3.5	6.2	0.6	3.3	2.7	1.0	2.6
4.3	4.0	2.9	4.9	1.2	3.4	3.0	0.9	2.3
5.1	5.3	4.5	6.1	1.4	3.1	2.9	1.1	2.7
4.7	3.1	5.8	6.6	0.9	2.0	2.5	0.9	2.0
4.3	2.9	3.8	5.7	1.0	2.0	2.6	0.7	2.3
5.3	4.7	5.1	6.0	2.0	3.0	2.6	1.1	2.6
5.3	4.3	4.9	6.5	1.2	3.0	2.3	1.4	2.8
3.1	2.6	5.0	5.2	0.6	2.1	2.5	1.0	2.4
4.4	6.7	2.2	6.4	2.2	3.0	1.1	1.7	3.0
3.9	4.8	3.3	4.5	1.1	3.7	2.7	1.0	2.6
4.8	4.8	4.3	5.9	1.0	3.4	2.7	1.1	2.8
4.3	5.2	4.2	6.6	2.5	3.0	1.9	2.2	2.8
5.3	5.1	5.0	6.1	1.5	3.2	2.5	1.4	2.8
5.2	4.1	4.7	5.7	1.0	3.0	2.6	1.4	2.7
4.3	4.7	1.6	5.4	1.3	3.0	1.1	1.1	2.9
5.1	4.7	4.3	6.0	1.3	3.0	2.8	1.3	2.6
4.9	4.9	4.0	5.3	1.7	3.0	2.4	1.8	2.8
4.3	5.1	3.4	5.1	2.0	3.1	1.9	1.6	2.8
4.7	4.0	4.9	5.7	2.5	3.2	2.4	1.1	2.4

5.5	4.7	4.6	6.4	1.8	3.0	2.4	1.4	2.8
4.9	4.1	3.6	5.6	1.2	3.1	2.6	1.3	2.6
5.9	5.2	5.3	7.0	2.1	3.1	2.9	1.3	2.6
4.4	3.7	4.2	6.1	1.0	2.9	2.4	1.4	2.7
5.1	4.1	4.4	6.3	1.8	3.1	2.5	1.5	2.6
5.7	5.9	4.1	6.0	1.0	3.1	2.5	1.2	2.9
4.9	3.9	3.4	4.7	2.7	3.0	1.9	1.7	2.9
4.3	4.1	4.0	6.2	0.7	3.0	2.5	1.7	2.8
5.2	4.1	4.4	6.2	0.9	3.0	2.2	1.4	2.9
4.8	3.8	3.8	5.2	1.6	3.0	2.4	2.1	2.8
5.1	4.8	2.4	4.6	2.2	3.0	1.4	2.0	3.0
4.8	3.8	3.6	5.9	1.5	3.1	2.6	0.9	2.8
5.6	5.6	3.4	5.6	1.9	3.0	1.6	1.4	3.0
5.6	4.8	3.5	5.9	2.0	3.1	2.1	1.3	2.9
4.7	4.0	4.1	6.6	1.1	3.0	2.9	1.5	2.8
5.2	4.8	4.3	6.9	1.5	3.2	2.5	0.8	2.7
4.9	4.5	4.0	6.4	0.8	3.0	2.7	1.6	2.9
4.4	5.1	5.7	6.4	2.0	3.2	2.8	1.5	2.7
4.6	5.1	2.3	6.2	2.5	3.0	1.7	1.3	2.9
4.7	3.8	1.6	5.9	5.1	3.0	1.0	0.9	2.7
4.7	3.6	2.1	6.9	4.8	3.0	1.0	0.9	2.9
5.6	3.3	1.8	5.9	2.5	3.0	1.0	0.9	3.0
5.4	5.2	5.1	6.3	1.4	3.0	2.1	2.0	3.0
-999	-999	-999	-999	-999	-999	-999	-999	-999
5.0	4.3	3.8	5.9	3.4	3.1	1.9	1.2	2.8
3.3	3.0	1.8	5.6	3.3	3.0	1.3	1.5	2.8
4.1	4.3	3.2	5.6	1.6	3.1	1.6	1.7	2.9
3.3	2.6	2.8	3.6	2.3	3.0	1.6	2.0	3.0
4.6	5.5	3.9	4.8	2.3	3.0	1.8	1.8	2.9
4.0	4.2	2.4	5.9	2.0	3.1	2.0	1.2	2.8
3.6	2.7	2.0	2.9	3.1	1.1	0.8	2.8	4.6
5.8	4.3	3.4	5.6	3.6	3.0	1.4	1.8	2.6
5.2	4.7	1.9	5.4	1.5	3.0	1.3	2.0	3.0
5.2	5.3	5.4	6.6	0.8	3.1	2.8	1.1	2.6
2.7	4.2	3.9	5.4	0.7	3.3	3.2	0.9	2.3
2.0	5.4	2.6	4.8	1.4	3.0	1.8	1.6	2.9
1.8	4.6	2.6	4.2	0.8	3.3	3.7	0.8	2.7
2.7	5.8	3.6	6.2	0.4	3.0	2.5	1.7	3.0
1.7	1.4	4.0	4.8	1.5	1.6	2.9	0.7	2.4
3.5	5.2	4.0	6.0	0.5	3.1	2.9	1.3	2.8
3.4	3.9	3.9	4.5	1.0	3.2	2.7	1.1	2.5
2.0	4.3	3.5	5.0	1.0	3.0	3.5	1.5	2.5
4.1	6.9	4.6	7.2	0.8	3.0	3.0	2.2	3.0
4.0	5.9	3.6	4.6	1.5	3.0	2.4	1.7	3.0
2.1	4.2	4.0	5.1	0.9	3.2	3.4	1.2	2.6
1.8	1.0	4.5	4.7	1.0	1.3	2.7	0.2	1.8
3.1	3.6	3.8	5.9	0.9	3.2	3.4	1.6	2.6
2.4	4.1	2.7	5.1	1.9	3.2	1.5	0.9	2.9
1.8	3.7	3.4	4.7	1.0	3.7	3.3	0.6	2.4
1.5	4.2	2.5	4.3	0.2	3.8	3.3	0.2	2.4
2.6	4.4	4.2	5.1	0.7	3.5	3.6	0.5	2.5
2.7	4.0	4.0	5.2	1.0	2.9	3.0	1.5	2.9
2.2	2.2	3.7	4.5	1.0	2.7	2.8	1.5	2.6
1.9	1.5	4.3	6.2	0.4	1.0	2.3	0.1	1.6
3.2	3.7	4.2	5.6	0.4	3.0	2.8	1.5	2.7
3.5	3.9	4.5	6.1	0.9	3.0	2.8	2.0	3.0

1.4	3.5	2.7	4.1	0.3	3.6	3.5	0.5	2.3
3.1	5.1	3.7	5.9	0.8	3.0	2.9	2.3	3.0
5.1	5.0	5.8	5.4	2.6	3.0	2.4	2.5	2.7
2.2	5.0	4.6	4.5	1.9	3.0	2.9	1.8	2.9
3.3	4.9	2.6	5.8	1.1	3.0	2.1	1.3	3.0
4.7	5.6	5.7	6.2	2.7	3.0	3.0	2.2	3.0
3.7	4.0	5.5	4.5	1.6	2.7	2.2	1.8	2.6
2.7	3.9	3.9	4.6	0.7	3.3	3.1	1.0	2.3
2.0	4.7	3.2	5.8	0.4	3.4	3.4	0.8	2.5
2.9	4.7	4.1	5.5	0.9	3.2	3.1	1.4	2.6
3.3	5.1	3.9	6.6	1.0	3.2	3.0	1.3	2.7
3.2	4.8	3.8	5.5	1.6	3.0	3.0	1.3	2.9
2.5	5.9	4.1	5.0	1.0	3.2	2.8	1.2	2.6
3.6	4.3	4.5	5.4	1.4	3.1	3.0	1.3	2.7
1.7	5.4	1.8	4.9	1.5	3.0	2.2	1.2	3.0
2.7	5.3	3.0	4.6	1.8	3.0	2.1	1.8	2.9
1.7	5.5	2.8	4.5	2.2	3.0	2.6	1.8	3.0
1.4	4.5	2.2	5.1	2.8	3.0	2.3	1.8	2.9
4.2	4.3	4.8	6.1	3.9	5.0	2.6	2.7	4.8
1.8	2.1	2.1	5.2	2.1	3.0	3.1	1.6	2.9
2.9	5.5	3.5	5.5	2.8	3.0	2.6	1.5	2.9
3.2	3.8	5.0	5.4	1.7	3.0	2.4	1.9	2.8
1.1	3.6	1.4	5.5	1.5	3.0	2.6	1.7	2.9
3.1	5.1	3.9	6.2	1.4	3.1	2.9	1.4	2.9
3.5	4.0	2.4	6.6	2.5	3.0	2.1	1.5	3.0
3.4	4.9	2.8	6.0	2.2	3.0	2.0	1.7	3.0
2.1	2.7	3.2	4.6	1.3	3.0	3.2	1.3	2.8
3.7	5.0	2.6	6.0	4.1	3.0	1.9	1.5	3.0
2.4	3.9	4.2	4.9	1.2	3.0	3.0	1.3	2.7
2.2	4.0	4.1	4.8	1.3	3.2	3.1	1.3	2.8
1.6	4.1	3.2	4.7	0.7	3.3	3.2	1.0	2.6
4.1	3.6	1.8	7.7	3.9	3.0	1.1	1.0	2.9
2.7	4.4	4.0	6.4	0.9	3.1	3.1	1.3	2.8
2.9	4.1	4.0	5.3	1.1	3.1	2.9	1.4	2.7
2.8	3.5	4.5	5.3	0.9	2.7	2.6	1.8	2.6
3.4	5.4	4.0	6.1	1.7	3.0	2.6	2.1	3.0
-999	4.4	5.0	7.1	1.4	2.9	2.6	1.3	3.0
-999	4.9	5.1	7.3	1.1	3.1	2.5	1.4	3.0
2.8	4.7	3.5	4.5	0.7	3.4	3.4	0.9	2.5
2.0	4.6	4.0	4.7	0.5	3.3	2.9	0.9	2.5
1.1	4.5	2.5	3.8	0.6	3.3	3.4	0.9	2.7
3.6	4.7	4.7	6.2	1.3	3.1	2.9	1.4	2.9
1.6	4.3	3.2	3.8	0.7	3.4	3.2	0.7	2.6
3.0	3.7	5.3	5.9	1.0	2.9	2.2	1.6	2.5
1.8	6.1	3.1	4.4	0.4	3.0	2.7	1.6	2.9

Legend

missing data: -999

sample with more than 1 external colour:

eg: 3+2 mixed 3 (orange) and 2: yellow, with

eg: 2+3 mixed 2 (yellow) and 3: orange, with

eg: 3(+2) mixed 3 (orange) and 2: yellow, wi

colour: in 2002: 99= pink salmon

Root skin colour	Root skin colour intensity	percentage external colour						Root cortex colour
		1. white	2. yellow	3. orange	4. red	5. purple	other	
3.0	5.1	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.5	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	5.0	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	5.0	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	5.0	0.000	0.000	1.000	0.000	0.000	0.000	3.0
5.0	3.8	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.9	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	5.0	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.8	0.000	0.000	1.000	0.000	0.000	0.000	3(+2)
3.0	5.0	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.0	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	5.0	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	5.0	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	5.0	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	5.3	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.8	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	5.0	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.8	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	5.3	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.6	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.9	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.2	0.000	0.000	1.000	0.000	0.000	0.000	3+2(+1)
3.0	4.8	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.9	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.6	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.9	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	5.0	0.000	0.000	1.000	0.000	0.000	0.000	3(+2)
3.0	5.0	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.8	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.9	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	5.0	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3+1+2	4.1	0.224	0.143	0.633	0.000	0.000	0.000	3+2+1
3.0	4.8	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	5.0	0.000	0.000	1.000	0.000	0.000	0.000	3.0
2+1+5	3.3	0.380	0.540	0.000	0.000	0.080	0.000	2+1
3.0	4.4	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.8	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3+1	3.3	0.340	0.000	0.660	0.000	0.000	0.000	3+1+2
3.0	4.5	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3+2+1(+5)	4.7	0.224	0.224	0.510	0.000	0.041	0.000	3+2+1
3+2+1(+5)	4.2	0.159	0.227	0.568	0.000	0.045	0.000	3+2+1
3(+2)	4.3	0.000	0.020	0.980	0.000	0.000	0.000	3(+2)

3(+2)	4.3	0.000	0.040	0.960	0.000	0.000	0.000	3(+2)
3.0	4.0	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.2	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.0	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.7	0.000	0.000	1.000	0.000	0.000	0.000	3(+1)
3+2(+1)	3.5	0.021	0.250	0.729	0.000	0.000	0.000	3+2(+1)
2+5+1+3	4.4	0.063	0.500	0.042	0.000	0.396	0.000	2+4+1
3.0	4.9	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.9	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3+2+1	4.4	0.042	0.167	0.792	0.000	0.000	0.000	3+2+1
1+3	4.0	0.700	0.000	0.300	0.000	0.000	0.000	1+2+3
3.0	5.0	0.000	0.000	1.000	0.000	0.000	0.000	3.0
2+1	5.5	0.360	0.640	0.000	0.000	0.000	0.000	2+1
5+2	5.7	0.174	0.087	0.000	0.000	0.739	0.000	2+4
3.0	4.8	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.7	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3+5	4.0	0.104	0.000	0.896	0.000	0.000	0.000	3.0
3+2+1	4.5	0.082	0.102	0.816	0.000	0.000	0.000	3+2(+1)
3+5+1+2	3.4	0.061	0.041	0.735	0.000	0.163	0.000	3+2+1+4
1.0	3.9	1.000	0.000	0.000	0.000	0.000	0.000	1+2
1.0	5.8	1.000	0.000	0.000	0.000	0.000	0.000	1.0
1.0	3.4	1.000	0.000	0.000	0.000	0.000	0.000	1.0
2+1	4.0	0.184	0.816	0.000	0.000	0.000	0.000	2+1
-999	-999	-999	-999	-999	-999	-999	-999	-999
3+2+1(+5)	4.0	0.260	0.300	0.400	0.000	0.040	0.000	2+3+1
1+2	4.5	0.857	0.143	0.000	0.000	0.000	0.000	1+2(+3)
5+2(+3)	4.2	0.000	0.040	0.020	0.000	0.940	0.000	2+1+4
5+1+3	2.5	0.080	0.000	0.040	0.000	0.880	0.000	1+2+3+4
3+1+2	3.5	0.080	0.040	0.880	0.000	0.000	0.000	3+2(+1)
3+1+2	4.6	0.300	0.180	0.520	0.000	0.000	0.000	3+2+1
3(+2+1)	2.9	0.020	0.040	0.940	0.000	0.000	0.000	3+2(+1)
5+1	5.5	0.040	0.000	0.000	0.000	0.960	0.000	2+4(+1)
5+1(+3)	4.3	0.400	0.000	0.020	0.000	0.580	0.000	1+2+4
3.0	4.9	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.0	0.000	0.000	1.000	0.000	0.000	0.000	3.0
2(+3)	5.0	0.000	0.980	0.020	0.000	0.000	0.000	2(+3)
3.0	3.9	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.7	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.8	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.9	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	3.5	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.8	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.4	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3(+2)	4.3	0.000	0.040	0.960	0.000	0.000	0.000	3(+2)
3.0	4.9	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.9	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.8	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	3.8	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.9	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.9	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.9	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	5.0	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.9	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	3.2	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.8	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.8	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	3.5	0.000	0.000	1.000	0.000	0.000	0.000	3.0

3.0	3.9	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	5.9	0.000	0.000	1.000	0.000	0.000	0.000	3.0
4+3	4.3	0.000	0.000	0.100	0.900	0.000	0.000	4+3+2
4.0	4.2	0.000	0.000	0.000	1.000	0.000	0.000	4.0
99+3+2+1*	3.8	0.160	0.060	0.380	0.000	0.000	0.400	99+3+2+1*
5.0	6.9	0.000	0.000	0.000	0.000	1.000	0.000	5(+2)
3.0	5.9	0.000	0.000	1.000	1.000	0.000	0.000	3.0
3.0	4.2	0.000	0.000	1.000	1.000	0.000	0.000	3.0
3.0	4.1	0.000	0.000	1.000	1.000	0.000	0.000	3.0
3.0	4.7	0.000	0.000	1.000	1.000	0.000	0.000	3.0
3.0	4.5	0.000	0.000	1.000	1.000	0.000	0.000	3.0
3.0	5.2	0.000	0.000	1.000	1.000	0.000	0.000	3.0
3+2	4.5	0.000	0.116	0.389	0.495	0.000	0.000	3+2
3.0	5.0	0.000	0.000	1.000	0.000	0.000	0.000	3.0
2+1+5+3	3.8	0.238	0.381	0.167	0.000	0.214	0.000	2+5+3
3+2+1(+5)	3.9	0.108	0.351	0.432	0.000	0.027	0.081	2+3
2+5+1+3	3.8	0.120	0.720	0.040	0.000	0.120	0.000	2+3+5
5+1+2	3.8	0.340	0.300	0.000	0.000	0.360	0.000	2+5
5+2(+3)	3.5	0.000	0.060	0.020	0.000	0.920	0.000	2(+3)
3.0	3.5	0.000	0.000	1.000	0.000	0.000	0.000	3.0
5+2+1	3.2	0.158	0.184	0.000	0.000	0.658	0.000	2+5
3+2	4.5	0.000	0.240	0.760	0.000	0.000	0.000	3+2
3+1	3.0	0.182	0.000	0.818	0.000	0.000	0.000	3+2
3+2	4.1	0.000	0.080	0.920	0.000	0.000	0.000	3+2
5+1+2	3.7	0.200	0.120	0.000	0.000	0.680	0.000	5+2+1
2+3+1	3.8	0.180	0.620	0.200	0.000	0.000	0.000	2+3+1
3.0	3.7	0.000	0.000	1.000	0.000	0.000	0.000	3.0
5+1+2+3	3.5	0.381	0.381	0.095	0.024	0.167	0.000	2+3(+1+5)
1.0	4.3	1.000	0.000	0.000	0.000	0.000	0.000	1.0
3.0	4.8	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	5.0	0.000	0.000	1.000	0.000	0.000	0.000	3.0
1+2+5	3.9	0.607	0.214	0.000	0.000	0.179	0.000	2+1
3.0	5.1	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.8	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.2	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	5.8	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.4	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	5.0	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.9	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.9	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.9	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	5.0	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	4.1	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	5.0	0.000	0.000	1.000	0.000	0.000	0.000	3.0
3.0	5.0	0.000	0.000	1.000	0.000	0.000	0.000	3.0

3 prevailing
 2 prevailing
 th 2 sporadic (<4%)

Root core colour	Percentage bolted
3.0	0.000
3.0	0.000
3+2	0.000
3+2	0.000
3.0	0.000
3+2	0.000
3+2	0.000
3+2	0.000
3+2	0.000
3.0	0.000
3(+2)	0.060
3.0	0.000
3.0	0.000
3.0	0.000
3.0	0.060
3+2	0.020
3+2	0.000
3+2	0.000
3.0	0.040
3+2	0.060
3.0	0.000
3+2(+1)	0.040
3.0	0.000
3.0	0.000
3+2	0.000
3+2	0.000
3(+2)	0.160
3.0	0.020
3.0	0.000
3.0	0.000
3(+2)	0.060
2+1(+3)	0.551
3.0	0.000
3.0	0.000
2+1	0.340
3+2	0.020
3+2	0.020
1+3+2	0.600
2+3	0.000
2+1+3	0.000
2+3+1	0.000
2+3	0.020

2+3	0.040
3+2	0.000
3.0	0.020
3+2	0.000
3(+1)	0.040
3+2+1	0.021
2+1	0.083
3(+2)	0.040
3.0	0.000
3+2+1	0.021
1+2	0.220
3.0	0.000
2+1	0.100
2.0	0.043
3(+2)	0.000
3(+2)	0.000
3(+2)	0.000
2(+3+1)	0.224
1+2+3	0.878
1+2	0.000
1(+2)	0.000
1.0	0.000
2+1	0.000
-999	-999
2+1+3	0.320
1+2	0.119
2+1+4(+3)	0.600
1+2(+3)	0.000
2+1+3	0.280
2+1+3	0.140
2+3(+1)	0.160
2(+1)	0.280
2+1	0.580
3.0	0.000
3.0	0.000
2(+3)	0.000
3(+2)	0.000
3+2	0.000
3.0	0.000
3(+2)	0.000
2+3	0.000
3.0	0.000
3+2	0.000
3+2	0.000
3.0	0.000
3.0	0.000
3+2	0.000
3.0	0.000
3+2	0.000
3(+2)	0.000
3.0	0.000
3.0	0.000
2+3	0.000
2+3	0.000
3(+2)	0.000
3+2	0.000

3.0	0.000
3.0	0.000
2+4+3+1	0.080
4+2+3	0.000
2+3+99+1*	0.080
5+2	0.000
3(+2)	0.000
2+3	0.000
3.0	0.000
3+2	0.000
3(+2)	0.000
3(+2)	0.000
3+2	0.000
3(+2)	0.000
2(+3)	0.000
2(+3)	0.027
2(+1)	0.000
2.0	0.000
2.0	0.000
3+2	0.000
2.0	0.000
3+2	0.000
3+2	0.000
2(+1)	0.420
2+1+3	0.120
3+2	0.000
2+1(+3)	0.571
3+2	0.000
3(+2)	0.000
3+2	0.000
2+1	0.321
3.0	0.000
3+2	0.000
3+2	0.000
3(+2)	0.000
3.0	0.000
3.0	0.000
3(+2)	0.000
3(+2)	0.000
3(+2)	0.000
3.0	0.000
3.0	0.000
3+2	0.000
3(+2)	0.000