## **Supplementary Material**

## Differentiation of metallicolous and non-metallicolous *Salix caprea* populations based on phenotypic characteristics and nuclear microsatellite (SSR) markers

Markus Puschenreiter<sup>1\*</sup>, Mine Türktaş<sup>2\*</sup>, Peter Sommer<sup>1</sup>, Gerlinde Wieshammer<sup>1</sup>, Gregor Laaha<sup>3</sup>, Walter W. Wenzel<sup>1</sup> and Marie-Theres Hauser<sup>2</sup>

**Supplementary Table 1:** Summary of the soil characteristics for each isolate and their growth and accumulation behaviour in perlite-based soil-less cultures exposed to elevated levels of Cd  $(0.5 \text{ mg L}^{-1})$  and Zn  $(5 \text{ mg L}^{-1})$ .

~ ''	-	<u></u>		Zn	Cd	Plant	Zn Perlite		Cd Perlite		Biomass	
Soil sample	Zn total	Cd total	pН	labile	labile	sample	a		1			
_	mg/kg	mg/kg		mg/kg	mg/kg	Perlite	mg/kg		mg/kg		(g)	
Prag						Nr	mean	SE	mean	SE	mean	SE
P1-5, 21-22 S	87.4	0.41	5.83	0.85	0.016	1	888	87.2	119	16.93	3.52	0.23
P1-5, 21-22 S	87.4	0.41	5.83	0.85	0.016	2	787	22.5	82.0	3.77	5.20	0.25
P1-5, 21-22 S	87.4	0.41	5.83	0.85	0.016	3	773	17.8	68.8	6.39	4.10	0.44
P1-5, 21-22 S	87.4	0.41	5.83	0.85	0.016	4	874	32.2	97.8	7.34	2.44	0.54
P1-5, 21-22 S	87.4	0.41	5.83	0.85	0.016	5	797	37.9	121	22.4	3.10	0.90
P6 S	67.9	0.26	6.06	0.37	0.008	6	638	10.0	72.6	2.57	2.99	0.44
P7-8 S a	72.0	0.28	7.62	0.12	0.008	7	600	10.1	58.5	4.08	5.04	0.24
P7-8 S b	72.0	0.28	7.62	0.12	0.005	8	577	8.8	46.5	1.76	5.29	1.15
P9 S	67.5	0.29	6.57	0.20	0.006	9	680	35.1	68.0	5.28	2.24	0.43
P10-11 S	67.5	0.26	6.48	0.17	0.005	10	1187	206	147	35.0	0.89	0.48
P10-11 S	67.5	0.26	6.48	0.17	0.005	11	922	51.3	91.6	1.54	6.06	0.40
P12-13 - S	59.9	0.19	5.43	0.65	0.016	12	799	65.8	90.8	9.42	4.20	1.10
P12-13 - S	59.9	0.19	5.43	0.65	0.016	13	777	41.0	85.3	7.11	5.78	1.02
P14-16+18S	64.6	0.22	5.43	0.80	0.019	14	678	21.6	71.5	7.22	4.35	0.51
P14-16+18S	64.6	0.22	5.43	0.80	0.019	15	835	94.5	94.0	13.9	4.22	0.34
P14-16+18S	64.6	0.22	5.43	0.80	0.019	16	913	44.0	141	31.1	2.53	0.53
P17 S	60.3	0.19	5.37	0.73	0.019	17	634	16.4	61.4	1.73	4.75	0.35
P14-16+18S	64.6	0.22	5.43	0.80	0.019	18	644	58.0	86.4	17.3	3.22	0.57
P1-5, 21-22 S	87.4	0.41	5.83	0.85	0.016	21	752	67.7	96.6	12.8	4.22	1.20
P1-5, 21-22 S	87.4	0.41	5.83	0.85	0.016	22	815	59.4	111	10.9	3.46	0.30
Mezica												
S - M1-3, 21-25	1292	18.1	7.2	0.92	0.10	1	820	19.6	119	15.4	3.24	0.28
S - M1-3, 21-25	1292	18.1	7.2	0.92	0.10	3	736	23.8	122	24.7	3.61	0.49
S - M5	9534	59.4	7.2	29.4	0.63	4	845	216	146	55.6	3.36	1.02
S - M7	404	7.65	7.27	0.70	0.07	7	608	39.6	86.4	2.9	5.35	0.21
S - M9+10+11	2747	13.6	7.57	2.35	0.04	10	784	114	90.0	3.1	3.07	0.85
S - M9+10+11	2747	13.6	7.57	2.35	0.04	11	1476	235	243	50.5	1.67	0.52
S - M8, 12-14, 26	1508	15.3	6.94	1.33	0.07	13	570	23.3	76.0	3.9	4.63	0.31
S - M15	3269	28.5	7.06	7.75	0.18	15	882	52.8	145	11.2	2.61	0.35
S - M16	489	9.94	7.19	1.05	0.06	16	679	35.7	112	9.0	2.63	0.35
S - M17	131	1.93	7.45	0.40	0.02	17	541	19.5	65.4	3.0	2.69	0.12
S - M18	2267	35.3	6.64	6.99	0.20	18	686	178	86.7	38.1	1.74	1.49
S - M19+20	1867	18.3	7.28	3.60	0.10	19	719	119	133	30.2	2.78	0.40

S - M1-3, 21-25       1292       18.1       7.2       0.92       0.10       21       1656       178       265       25.3       1.64       0.28         S - M1-3, 21-25       1292       18.1       7.2       0.92       0.10       22       893       47.4       112       9.0       4.13       0.32         S - M1-3, 21-25       1292       18.1       7.2       0.92       0.10       23       995       149       156       26.0       2.68       0.56         S - M1-3, 21-25       1292       18.1       7.2       0.92       0.10       24       1228       51.4       207       16.3       1.55       0.17         S - M1-3, 21-25       1292       18.1       7.2       0.92       0.10       24       1228       51.4       207       16.3       1.55       0.17         S - M1-3, 21-25       1292       18.1       7.2       0.92       0.10       25       1103       120       152       12.1       2.60       0.66         S - M8, 12-14, 26       1508       15.3       6.94       1.33       0.07       26       959       64.3       140       9.4       2.89       0.29         S - M18, 27
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S - M1-3, 21-25       1292       18.1       7.2       0.92       0.10       23       995       149       156       26.0       2.68       0.56         S - M1-3, 21-25       1292       18.1       7.2       0.92       0.10       24       1228       51.4       207       16.3       1.55       0.17         S - M1-3, 21-25       1292       18.1       7.2       0.92       0.10       25       1103       120       152       12.1       2.60       0.66         S - M3, 21-25       1292       18.1       7.2       0.92       0.10       25       1103       120       152       12.1       2.60       0.66         S - M8, 12-14, 26       1508       15.3       6.94       1.33       0.07       26       959       64.3       140       9.4       2.89       0.29         S - M18, 27       2267       35.3       6.64       6.99       0.20       27       1154       212       138       9.2       1.87       0.82         Pribram
S - M1-3, 21-25       1292       18.1       7.2       0.92       0.10       24       1228       51.4       207       16.3       1.55       0.17         S - M1-3, 21-25       1292       18.1       7.2       0.92       0.10       25       1103       120       152       12.1       2.60       0.66         S - M8, 12-14, 26       1508       15.3       6.94       1.33       0.07       26       959       64.3       140       9.4       2.89       0.29         S - M18, 27       2267       35.3       6.64       6.99       0.20       27       1154       212       138       9.2       1.87       0.82         S - M18, 27       2267       35.3       6.64       6.99       0.20       27       1154       212       138       9.2       1.87       0.82         Delta       28       781       38.9       104       11.6       2.56       0.23         Pribram       276       7.35       4.88       25.7       1.68       1       765       46.7       182       29.1       1.01       0.08         PR 2-6       202       4.70       5.5       13.1       1.22       2       743
S - M1-3, 21-25       1292       18.1       7.2       0.92       0.10       25       1103       120       152       12.1       2.60       0.66         S - M8, 12-14, 26       1508       15.3       6.94       1.33       0.07       26       959       64.3       140       9.4       2.89       0.29         S - M18, 27       2267       35.3       6.64       6.99       0.20       27       1154       212       138       9.2       1.87       0.82         S - M18, 27       2267       35.3       6.64       6.99       0.20       27       1154       212       138       9.2       1.87       0.82         Pribram
S - M8, 12-14, 26       1508       15.3       6.94       1.33       0.07       26       959       64.3       140       9.4       2.89       0.29         S - M18, 27       2267       35.3       6.64       6.99       0.20       27       1154       212       138       9.2       1.87       0.82         Pribram       28       781       38.9       104       11.6       2.56       0.23         PR1 S       276       7.35       4.88       25.7       1.68       1       765       46.7       182       29.1       1.01       0.08         PR 2-6       202       4.70       5.5       13.1       1.22       2       743       28.4       90.8       6.51       1.74       0.26         PR 2-6       202       4.70       5.5       13.1       1.22       3       751       45.1       85.4       6.35       2.91       0.32
S - M18, 27       2267       35.3       6.64       6.99       0.20       27       1154       212       138       9.2       1.87       0.82         Pribram       28       781       38.9       104       11.6       2.56       0.23         PR1 S       276       7.35       4.88       25.7       1.68       1       765       46.7       182       29.1       1.01       0.08         PR 2-6       202       4.70       5.5       13.1       1.22       2       743       28.4       90.8       6.51       1.74       0.26         PR 2-6       202       4.70       5.5       13.1       1.22       3       751       45.1       85.4       6.35       2.91       0.32
Pribram         28         781         38.9         104         11.6         2.56         0.23           Pribram         276         7.35         4.88         25.7         1.68         1         765         46.7         182         29.1         1.01         0.08           PR 2-6         202         4.70         5.5         13.1         1.22         2         743         28.4         90.8         6.51         1.74         0.26           PR 2-6         202         4.70         5.5         13.1         1.22         3         751         45.1         85.4         6.35         2.91         0.32
Pribram         Image: Constraint of the state of t
PR1 S         276         7.35         4.88         25.7         1.68         1         765         46.7         182         29.1         1.01         0.08           PR 2-6         202         4.70         5.5         13.1         1.22         2         743         28.4         90.8         6.51         1.74         0.26           PR 2-6         202         4.70         5.5         13.1         1.22         3         751         45.1         85.4         6.35         2.91         0.32
PR 2-6         202         4.70         5.5         13.1         1.22         2         743         28.4         90.8         6.51         1.74         0.26           PR 2-6         202         4.70         5.5         13.1         1.22         3         751         45.1         85.4         6.35         2.91         0.32
PR 2-6 202 4.70 5.5 13.1 1.22 3 751 45.1 85.4 6.35 2.91 0.32
PR 2-6 202 4.70 5.5 13.1 1.22 4 929 63.8 111 16.8 1.74 0.65
PR 2-6 202 4.70 5.5 13.1 1.22 5 722 64.4 65.3 21.6 2.80 0.55
PR 2-6 202 4.70 5.5 13.1 1.22 6 614 33.4 52.9 2.88 3.21 0.28
PR7-9 S 4182 35.8 6.35 36.5 1.01 7 936 26.6 137 0.56 1.40 0.35
PR7-9 S 4182 35.8 6.35 36.5 1.01 8 648 47.7 67.7 10.8 3.59 0.47
PR7-9 S 4182 35.8 6.35 36.5 1.01 9 670 38.7 84.1 10.5 2.50 0.25
PR10-12 S a 4387 38.9 6.5 63.3 1.40 10 928 20.9 76.5 8.26 1.55 0.18
PR10-12 S a 4387 38.9 6.5 63.3 1.40 11 807 63.0 141 9.29 1.25 0.18
PR13 S 1139 10.3 7.35 2.01 0.13 13 715 55.5 65.2 8.44 2.74 0.33
PR14-17 S 2091 12.6 7.25 1.20 0.11 14 838 90.8 134 52.6 1.65 0.52
PR14-17 S 2091 12.6 7.25 1.20 0.11 15 859 57.3 87.6 3.84 2.00 0.12
PR14-17 S 2091 12.6 7.25 1.20 0.11 16 624 47.2 67.4 6.97 2.78 0.16
PR18-19 S 2962 30.0 5.95 221 4.69 18 879 67.8 129 14.8 1.98 0.11
PR18-19 S 2962 30.0 5.95 221 4.69 19 774 75.9 86.9 12.2 1.78 0.09
Arnoldstein
A1-3, A21-S 2492 19.4 7.3 14.3 0.11 1 913 27.0 148 13.8 1.85 0.24
A5, A22-S 1530 11.9 6.23 57.4 0.90 5 701 22.5 118 8.61 3.00 0.38
A 6 - 8-S 1244 9.65 6.61 8.68 0.21 6 685 44.2 88.6 12.1 5.09 0.40
A 6 - 8-S 1244 9.65 6.61 8.68 0.21 8 709 68.7 138 25.7 2.13 0.40
A9-S 4465 13.5 7.89 35.1 0.17 9 590 45.9 79.0 7.97 3.84 0.22
A10-16-S 1824 3.91 7.97 7.57 0.03 13 604 67.4 100 8.38 3.51 0.30
A10-16-S 1824 3.91 7.97 7.57 0.03 11 545 23.8 78.6 1.64 3.37 0.09
A10-16-S 1824 3.91 7.97 7.57 0.03 14 733 68.0 117 15.6 3.20 0.42
A10-16-S 1824 3.91 7.97 7.57 0.03 15 900 85.9 192 25.0 2.34 0.37
A17-S 1011 3.93 7.59 0.82 0.02 16 784 25.0 129 3.55 2.87 0.61
A18-S 1898 12.0 7.44 10.0 0.15 19 903 130 153 30.5 2.82 0.51
A19,20-S 1134 6.50 7.49 2.83 0.10 20 516 57.0 65.3 16.8 4.56 0.89
A1-3, A21-S 2492 19.4 7.3 14.3 0.11 21 855 100 127 20.0 1.85 0.70
A5, A22-S 1530 11.9 6.23 57.4 0.90 22 1062 90.2 169 27.1 3.13 0.63
A19,20-S=A23 1134 6.50 7.49 2.83 0.10 23 751 36.5 122 7.68 2.47 0.09
A1-3, 21, 24 S 2492 19.4 7.3 14.3 0.11 24 1192 57.0 248 15.1 1.52 0.10
A18-S=A27 1898 12.0 7.44 10.0 0.15 27 1173 28.0 192 28.1 2.17 0.31
A9-S=A28 4465 13.5 7.89 35.1 0.17 28 910 108 179 23.6 1.16 0.18
Forchtenstein
F2-4S 80.9 0.46 5.83 5.94 0.052 2 691 79.6 126 19.5 2.24 0.53
F2-4S 80.9 0.46 5.83 5.94 0.052 3 606 32.4 72.1 3.13 3.23 0.31
F5-9S 92.8 0.37 4.74 3.50 0.032 6 681 65.7 83.1 12.7 4.10 0.56
F5-9S 92.8 0.37 4.74 3.50 0.032 7 570 21.7 60.1 4.28 4.40 0.19
F5-9S 92.8 0.37 4.74 3.50 0.032 8 917 98.3 114 19.1 4.04 0.80
F10-18S         89.0         0.59         4.3         8.03         0.049         10         804         47.4         96.6         16.6         2.90         0.52

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F10-18S       89.0       0.59       4.3       8.03       0.049       13       659       26.2       93.4       3.32       3.45       0.23         F10-18S       89.0       0.59       4.3       8.03       0.049       14       797       110       125       26.4       2.75       0.41         F10-18S       89.0       0.59       4.3       8.03       0.049       17       633       11.5       85.1       2.85       3.74       0.49         F10-18S       519*       0.50       5.14       76.45       0.028       19       632       37.5       61.8       4.76       3.20       0.29         F20-S       228*       0.79       7.47       0.15       0.003       20       575       48.4       57.5       9.65       4.37       1.00         Kutna Hora
F10-18S89.00.594.38.030.0491479711012526.42.750.41F10-18S89.00.594.38.030.0491763311.585.12.853.740.49F19-S519*0.505.1476.450.0281963237.561.84.763.200.29F20-S228*0.797.470.150.0032057548.457.59.654.371.00Kutna Hora </td
F10-18S       89.0       0.59       4.3       8.03       0.049       17       633       11.5       85.1       2.85       3.74       0.49         F19-S       519*       0.50       5.14       76.45       0.028       19       632       37.5       61.8       4.76       3.20       0.29         F20-S       228*       0.79       7.47       0.15       0.003       20       575       48.4       57.5       9.65       4.37       1.00         Kutna Hora   <
F19-S       519*       0.50       5.14       76.45       0.028       19       632       37.5       61.8       4.76       3.20       0.29         F20-S       228*       0.79       7.47       0.15       0.003       20       575       48.4       57.5       9.65       4.37       1.00         Kutna Hora                          61.8       4.76       3.20       0.29          Kutna Hora
F20-S       228*       0.79       7.47       0.15       0.003       20       575       48.4       57.5       9.65       4.37       1.00         Kutna Hora <td< td=""></td<>
Kutna HoraImage: Constraint of the state of t
KH1-2 S860372.006.4282.991.206192429.21339.551.740.38KH1-2 S860372.006.4282.991.206274574.176.26.962.400.34KH3-5, 21 S274031.507.915.360.124474154.310415.02.570.56KH3-5, 21 S274031.507.915.360.124565019.961.04.882.370.49KH6 S822084.007.1414.650.378686261.511017.01.660.36KH7 S6304.007.581.940.024792159.911711.32.340.25KH8 S501554.007.199.360.364866727.682.22.921.790.15KH10, 22 S481461.007.0218.520.6731061833.147.35.423.910.46KH11-12S401056.007.0816.190.3531195518585.29.611.630.22KH13 S5054.508.341.060.0051376715284.516.02.270.26KH14 S5154.007.072.920.1411461738.988.67.132.380.14KH15, 23 S180114.005.121
KH1-2 S860372.006.4282.991.206274574.176.26.962.400.34KH3-5, 21 S274031.507.915.360.124474154.310415.02.570.56KH3-5, 21 S274031.507.915.360.124565019.961.04.882.370.49KH6 S822084.007.1414.650.378686261.511017.01.660.36KH7 S6304.007.581.940.024792159.911711.32.340.25KH8 S501554.007.199.360.364866727.682.22.921.790.15KH10, 22 S481461.007.0218.520.6731061833.147.35.423.910.46KH11-12S401056.007.0816.190.3531195518585.29.611.630.22KH13 S5054.508.341.060.0051376715284.516.02.270.26KH14 S5154.007.072.920.1411461738.988.67.132.380.14KH15, 23 S180114.005.1212.740.2721593172.211926.91.210.19KH16, 17 S7154.007.52 <td< td=""></td<>
KH3-5, 21 S274031.507.915.360.124474154.310415.02.570.56KH3-5, 21 S274031.507.915.360.124565019.961.04.882.370.49KH6 S822084.007.1414.650.378686261.511017.01.660.36KH7 S6304.007.581.940.024792159.911711.32.340.25KH8 S501554.007.199.360.364866727.682.22.921.790.15KH10, 22 S481461.007.0218.520.6731061833.147.35.423.910.46KH11-12S401056.007.0816.190.3531195518585.29.611.630.22KH13 S5054.508.341.060.0051376715284.516.02.270.26KH14 S5154.007.072.920.1411461738.988.67.132.380.14KH15, 23 S180114.005.1212.740.2721593172.211926.91.210.19KH16, 17 S7154.007.54.550.1001690762.012414.60.25
KH3-5, 21 S274031.507.915.360.124565019.961.04.882.370.49KH6 S822084.007.1414.650.378686261.511017.01.660.36KH7 S6304.007.581.940.024792159.911711.32.340.25KH8 S501554.007.199.360.364866727.682.22.921.790.15KH10, 22 S481461.007.0218.520.6731061833.147.35.423.910.46KH11-12S401056.007.0816.190.3531195518585.29.611.630.22KH13 S5054.508.341.060.0051376715284.516.02.270.26KH14 S5154.007.072.920.1411461738.988.67.132.380.14KH15, 23 S180114.005.1212.740.2721593172.211926.91.210.19
KH6 S822084.007.1414.650.378686261.511017.01.660.36KH7 S6304.007.581.940.024792159.911711.32.340.25KH8 S501554.007.199.360.364866727.682.22.921.790.15KH10, 22 S481461.007.0218.520.6731061833.147.35.423.910.46KH11-12S401056.007.0816.190.3531195518585.29.611.630.22KH13 S5054.508.341.060.0051376715284.516.02.270.26KH14 S5154.007.072.920.1411461738.988.67.132.380.14KH15, 23 S180114.005.1212.740.2721593172.211926.91.210.19
KH7 S6304.007.581.940.024792159.911711.32.340.25KH8 S501554.007.199.360.364866727.682.22.921.790.15KH10, 22 S481461.007.0218.520.6731061833.147.35.423.910.46KH11-12S401056.007.0816.190.3531195518585.29.611.630.22KH13 S5054.508.341.060.0051376715284.516.02.270.26KH14 S5154.007.072.920.1411461738.988.67.132.380.14KH15, 23 S180114.005.1212.740.2721593172.211926.91.210.19KH16, 17 S7154.007.54.550.1201620762.012414.62.40
KH8 S501554.007.199.360.364866727.682.22.921.790.15KH10, 22 S481461.007.0218.520.6731061833.147.35.423.910.46KH11-12S401056.007.0816.190.3531195518585.29.611.630.22KH13 S5054.508.341.060.0051376715284.516.02.270.26KH14 S5154.007.072.920.1411461738.988.67.132.380.14KH15, 23 S180114.005.1212.740.2721593172.211926.91.210.19KH14 (17 S)7154.097.54.550.1901.690762.012414.62.400.25
KH10, 22 S       4814       61.00       7.02       18.52       0.673       10       618       33.1       47.3       5.42       3.91       0.46         KH11-12S       4010       56.00       7.08       16.19       0.353       11       955       185       85.2       9.61       1.63       0.22         KH13 S       505       4.50       8.34       1.06       0.005       13       767       152       84.5       16.0       2.27       0.26         KH14 S       515       4.00       7.07       2.92       0.141       14       617       38.9       88.6       7.13       2.38       0.14         KH15, 23 S       1801       14.00       5.12       12.74       0.272       15       931       72.2       119       26.9       1.21       0.19         KH16, 17 S       715       4.00       7.7       4.55       0.192       16       937       630       124       14.6       2.45
KH11-12S         4010         56.00         7.08         16.19         0.353         11         955         185         85.2         9.61         1.63         0.22           KH13 S         505         4.50         8.34         1.06         0.005         13         767         152         84.5         16.0         2.27         0.26           KH14 S         515         4.00         7.07         2.92         0.141         14         617         38.9         88.6         7.13         2.38         0.14           KH15, 23 S         1801         14.00         5.12         12.74         0.272         15         931         72.2         119         26.9         1.21         0.19           KH16, 17 S         715         4.55         0.192         16         907         62.0         124         14.6         2.45
KH13 S         505         4.50         8.34         1.06         0.005         13         767         152         84.5         16.0         2.27         0.26           KH14 S         515         4.00         7.07         2.92         0.141         14         617         38.9         88.6         7.13         2.38         0.14           KH15, 23 S         1801         14.00         5.12         12.74         0.272         15         931         72.2         119         26.9         1.21         0.19
KH14 S         515         4.00         7.07         2.92         0.141         14         617         38.9         88.6         7.13         2.38         0.14           KH15, 23 S         1801         14.00         5.12         12.74         0.272         15         931         72.2         119         26.9         1.21         0.19           KH16, 17 S         715         4.00         7.5         4.55         0.120         16         207         62.0         124         14.6         2.45         0.25
KH15, 23 S         1801         14.00         5.12         12.74         0.272         15         931         72.2         119         26.9         1.21         0.19           KH16, 17 S         715         4.00         75         4.55         0.100         16         0.07         (2.0)         124         14.6         2.40         0.25
KH10-1/5 /15 4.00 /.5 4.55 0.180 16 80/ 63.0 124 14.6 2.49 0.35
KH16-17 S         715         4.00         7.5         4.55         0.180         17         699         46.4         68.2         9.00         1.54         0.30
KH11-12S         4010         56.00         7.08         16.19         0.353         12         859         70.9         105         7.32         4.10         0.28
KH18 S         109         1.50         7.67         1.685         0.045         18         768         107         123         21.8         2.81         0.33
KH19-20         105         1.50         8.46         0.88         0.005         19         745         96.2         86.1         14.5         2.67         0.39
KH19-20         105         1.50         8.46         0.88         0.005         20         667         21.8         116         7.73         2.26         0.28
KH3-5, 21 S         2740         31.50         7.91         5.36         0.124         21         1021         31.1         252         12.5         1.44         0.13
KH10, 22 S         4814         61.00         7.02         18.52         0.673         22         996         59.7         215         35.6         1.68         0.55
KH15, 23 S         1801         14.00         5.12         12.74         0.272         23         1044         62.0         178         12.1         2.79         0.26
Völkermarkt
V1-5 63.5 0.58 6.8 0.58 0.002 1 1042 296 172 78.9 2.60 0.87
V1-5 63.5 0.58 6.8 0.58 0.002 2 927 72.8 142 23.0 2.34 0.48
V1-5 63.5 0.58 6.8 0.58 0.002 3 963 33.5 145 8.70 2.67 0.21
V1-5 63.5 0.58 6.8 0.58 0.002 4 868 170 160 27.4 2.43 0.11
V1-5 63.5 0.58 6.8 0.58 0.002 5 1116 261 190 28.6 1.14 0.4/
V6-11         /4.4         0.46         6.7         0.72         0.010         7         925         158         154         34.8         2.56         0.64           V6-11         74.4         0.46         6.7         0.72         0.010         7         925         158         154         34.8         2.56         0.64           V6-11         74.4         0.46         6.7         0.72         0.010         0         664         94.6         102         20.7         2.05         0.41
V6-11 $/4.4$ $0.46$ $6.7$ $0.72$ $0.010$ $9$ $664$ $84.6$ $103$ $20.7$ $3.05$ $0.41$ $V6-11$ $74.4$ $0.46$ $6.7$ $0.72$ $0.010$ $10$ $020$ $76.1$ $147$ $16.8$ $2.62$ $0.26$
V6-11         /4.4         0.40         6.7         0.72         0.010         10         939         /6.1         147         16.8         2.63         0.20           V6-11         74.4         0.46         6.7         0.72         0.010         11         939         /6.1         147         16.8         2.63         0.20           V6-11         74.4         0.46         6.7         0.72         0.010         11         989         64.7         150         6.21         0.82         0.14
V0-11 $/4.4$ $0.40$ $0.7$ $0.72$ $0.010$ $11$ $888$ $04.7$ $159$ $0.21$ $0.82$ $0.14$ $V12$ 21         58.4 $0.45$ 6.5 $1.40$ $0.001$ $12$ $704$ $22.7$ $117$ $0.10$ $2.55$ $0.31$
$v_{12-21}$ $38.4$ $0.43$ $0.5$ $1.40$ $0.001$ $12$ $/94$ $22.7$ $117$ $9.19$ $2.55$ $0.51$ $v_{12}$ $22$ $58.4$ $0.45$ $6.5$ $1.40$ $0.001$ $12$ $/94$ $22.7$ $117$ $9.19$ $2.55$ $0.51$ $v_{12}$ $22$ $58.4$ $0.45$ $6.5$ $1.40$ $0.001$ $12$ $994$ $22.7$ $117$ $9.19$ $2.55$ $0.51$
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$
V12-28 58.4 0.45 6.5 1.40 0.001 19 1064 192 160 28.8 1.87 0.75
V12-29 58.4 0.45 6.5 1.40 0.001 20 759 59.0 106 16.0 3.95 0.41
V12-30 58.4 0.45 6.5 1.40 0.001 21 1009 150 109 12.7 2.23 0.75

\*) these two Zn soil concentrations are probably high because of a nearby rusty and leaching fence.

Locus	Allele Size	Primer Sequence 5` - 3`	Motif in <i>S. caprea</i>	Motif in P. trichocarpa	Chromsomal Position
ORPM_62	170-198	F: CGGAGTCAGCTTGAGGTAGC	[AT] <sub>2</sub> [T] <sub>8</sub>	[AT] <sub>4</sub> [ATTTT] <sub>3</sub>	Ι
		R: CGGCAATATTGAGGAGAATGA			16.99 Mb
ORPM_312	177-207	F: GTGGGGATCAATCCAAAAGA	[CTT] <sub>5</sub>	[CCT] <sub>6</sub>	VII
		R: CCCATATCAAACCATTTGAAAAA			9.31 Mb
ORPM_446	220-258	F: GGGCTGCAGACAAATTAAGG	[CT] <sub>3</sub> [CT] <sub>4</sub>	[CT] <sub>3</sub> [CT] <sub>4</sub>	XIV – 3.69
		R: TGGGACATGCTCCATGGTAT			Mb
				Motif in	
				P. nigra	
WMPS_12	146-155	F: TTTTTCGTATTCTTATCTATCC	[CA] <sub>4</sub>	[CA] <sub>4</sub>	VI
		R: CACTACTCTGACAAAACCATC			11.36 Mb
WMPS_14	205-237	F: CAGCCGCAGCCACTGAGAAATC	<sup>1</sup> [CAG] <sub>21</sub>	$^{1}[CAG]_{28}$	V
		R: GCCTGCTGAGAAGACTGCCTTGAC			12.53 Mb
WMPS_19	152-209	F: AGCCACAGCAAATTCAGATGATGC	<sup>2</sup> [CAG] <sub>22</sub>	<sup>2</sup> [CAG] <sub>38</sub>	V
		R: CCTGCTGAGAAGACTGCCTTGACA			12.53 Mb
WMPS_21	157-184	F: TGCTGATGCAAAAGATTTAG	[GCT] <sub>27</sub>	[GCT] <sub>45</sub>	II
		R: TTGGAACTTCAACATTCAGAT			5.05 Mb
				Motif in	
				S. burjatica	
SB_24	127-153	F: ACTTCAATCTCTCTGTATTCT	<sup>3</sup> [TG] <sub>9</sub>	<sup>3</sup> [TG] <sub>9</sub>	XI
		R: CTATTTATGGGTTGGTCGATC			4.70 Mb
SB_38	91-151	F: CCACTTGAGGAGTGTAAGGAT	[TG] <sub>27</sub>	[TG] <sub>27</sub>	IX
		R: CTTAAATGTAAAACTGAATCT			10.38 Mb
SB_199	95-127	F: CTATTTGGTCTCAATCACCTT	[TG] <sub>15</sub>	[TG] <sub>11</sub> CG[TG] <sub>6</sub>	XV
		R: CTTTACCTCAGAAAATCCAGA			9.74 Mb
				Motif in	
				S. lanata	
gSIMCT024	96-124	F: CTCCCTTCACTTGCTCCAT	[CT] <sub>14</sub>	[CT] <sub>10</sub>	Х
		R: TAATACCAGCCCTTAAAGAAG			14.27 Mb

## Supplementary Table 2 Overview of the SSR marker characteristics

<sup>1</sup>Complete complex motif in *S. caprea*: [CAG]<sub>4</sub>CAT[CAG]<sub>12</sub>CAT[CAG]<sub>3</sub> and in *P. nigra*: [CAG]<sub>1</sub>CAT[CAG]<sub>12</sub>CAT[CAG]<sub>9</sub>CAT[CAG]<sub>3</sub>

<sup>2</sup>Complete complex motif in *S. caprea*: [CAG]<sub>2</sub>CAT[CAG]<sub>8</sub>CAT[CAG]<sub>10</sub> and in *P. nigra*: [CAG]<sub>3</sub>CAT[CAG]<sub>8</sub>CAT[CAG]<sub>8</sub>CAT[CAG]<sub>4</sub>CAA[CAG]<sub>3</sub>

<sup>3</sup>Complete complex motif in *S. caprea*: [TG]<sub>9</sub>A[TG]<sub>2</sub>A[TG]<sub>4</sub> and in *S. burjatica*: [TG]<sub>9</sub>A[TG]<sub>2</sub>A[TG]<sub>4</sub>...[TG]<sub>3</sub>

Supplementary Table 3 Heterozygosities, lnRH and normalized lnRH values of the 11 loci and seven populations. Loci with Norm. $\Theta$  values of larger +1.95 and smaller -1.95 have a significant reduced variability and thus might have been under selection and contribute to adaptation in the contamination area.

Locus	$H_{\rm obs}$	$H_{\rm exp}$	Populations							$\mu_{c}$	$\mu_{\mathrm{u}}$	lnRH	Norm.Θ
			A	М	V	F	KH	PR	Р			Θ	
ORPM_62	0.49	0.74	0.79	0.55	0.67	0.70	0.75	0.78	0.71	0,72	0,69	0,18	-0,30
ORPM_312	0.81	0.82	0.85	0.82	0.84	0.73	0.88	0.77	0.74	0,83	0,77	0,63	0,79
ORPM_446	0.56	0.81	0.86	0.80	0.79	0.75	0.80	0.79	0.80	0,81	0,78	0,33	0,07
WMPS_12	0.34	0.33	0.25	0.44	0.09	0.38	0.41	0.50	0.20	0,40	0,22	0,99	1,68
WMPS_14	0.85	0.84	0.81	0.86	0.84	0.85	0.77	0.86	0.90	0,83	0,86	-0,51	-1,97*
WMPS_19	0.84	0.88	0.86	0.88	0.85	0.86	0.88	0.89	0.89	0,88	0,87	0,17	-0,32
WMPS_21	0.79	0.76	0.75	0.75	0.75	0.71	0.75	0.76	0.82	0,75	0,76	-0,07	-0,90
SB_24	0.89	0.77	0.77	0.82	0.79	0.79	0.79	0.78	0.67	0,79	0,75	0,37	0,16
SB_38	0.82	0.91	0.90	0.93	0.82	0.90	0.92	0.89	0.89	0,91	0,87	0,74	1,07
SB_199	0.61	0.76	0.80	0.79	0.79	0.81	0.78	0.65	0.65	0,76	0,75	0,04	-0,63
gSIMCT024	0.85	0.80	0.82	0.74	0.73	0.84	0.84	0.83	0.76	0,82	0,78	0,45	0,35
all loci	0.71	0.77	0.77	0.76	0.72	0.76	0.78	0.77	0.73				

\* only significant without Bonferroni correction



**Supplementary Figure 1:** Shoot length (A) and leaf number (B) of *S. caprea* isolates. Boxes represent the median (vertical solid line), the arithmetic mean (vertical dashed line), and 25 - 75 % percentile. Whiskers represent the 90<sup>th</sup> and 10<sup>th</sup> percentile. Significant differences were determined by a post hoc comparison of means (Scheffé test after nested analysis of variance; p < 0.05) and are indicated by different letters.



**Supplementary Figure 2:** Geographic map indicating the location of the *S. caprea* populations. South the Alps in Slovenia the contaminated population in Mežica – M, and in Austria the contaminated population Arnoldstein – (A) and the non-contaminated sites, Völkermarkt - V and Forchtenstein – F. North the Alps in the Czech Republic the two contaminated populations Příbram – PR and Kutná Hora – KH and the non-contaminated site near Prague (P).



**Supplementary Figure 3:** The graphs show that the amount of labile Zn and Cd depends with a significance level for Pearson's correlation of p<0.01 on the total Zn and Cd concentration in the soil. In fact 40% of the variation of labile Zn and 36% for Cd are explained by the total Zn and Cd concentration, respectively. Other factors influencing the labile heavy metal fractions are pH, content of clay, carbonate and organic matter.



**Supplementary Figure 4:** Pearson's correlations between the level of contamination where the isolate originated and the biomass production in perlite cultures exposed to elevated levels of Cd and Zn. A and B are the graphs for the uncontaminated, C and D for the contaminated sites. The only significant negative correlation between Cd concentration and biomass production was seen at uncontaminated sites (A). Isolates from the contaminated sites did not show such a correlation indicating that they might have been selected to withstand higher Cd and Zn concentrations.

Note that two soil samples from the uncontaminated sites had very high Zn concentrations probably because of a nearby rusty fence that leached into the soil (B).



**Supplementary Figure 5:** Pearson's correlations between the level of soil contamination and Cd concentration in leaves after the exposure of the isolates in perlite to Cd and Zn. While a significant trend was found between soil contamination and accumulation capacity in perlite cultures for soil contamination below 20 mg kg<sup>-1</sup> (A, B) above this soil Cd contamination level the trend diminished.