

Table S1. 38 stress-related vegetation indices in the literature used in the study for SVM modeling.

No.	Name	Formula	Reference*
1	Carter	R_{695} / R_{420}	[1]
2	Carter2	R_{695} / R_{760}	[1]
3	Carter3	R_{605} / R_{760}	[1]
4	Carter4	R_{710} / R_{760}	[1]
5	Carter5	R_{695} / R_{670}	[1]
6	Carter6	R_{550}	[1]
7	CI	$R_{675} * R_{690} / R^2_{683}$	[2]
8	D730/D706	D_{730} / D_{706}	[2]
9	DDI	$(R_{720+\Delta} - R_{720}) - (R_{672+\Delta} - R_{672})$	[3]
10	DPI	$(D_{688} * D_{710}) / D_{697}^2$	[2]
11	DWSI2	R_{800} / R_{1660}	[4]
12	DWSI	R_{1660} / R_{550}	[4]
13	DWSI3	R_{1660} / R_{680}	[4]
14	DWSI4	R_{550} / R_{680}	[4]
15	DWSI5	$(R_{800} + R_{550}) / (R_{1660} + R_{680})$	[4]
16	EGFN	$\frac{(max(D_{650:750}) - max(D_{500:550}))}{(max(D_{650:750}) + max(D_{500:550}))}$	[5]
17	EGFR	$max(D_{650:750}) / max(D_{500:550})$	[5]
18	MPRI	$(R_{515} - R_{530}) / (R_{515} + R_{530})$	[6]
19	MCARI	$((R_{700} - R_{670}) - 0.2 * (R_{700} - R_{550})) * (R_{700} / R_{670})$	[7]
20	MSI	R_{1600} / R_{817}	[8]
21	MTCI	$(R_{754} - R_{709}) / (R_{709} - R_{681})$	[9]
22	NDVI	$(R_{800} - R_{680}) / (R_{800} + R_{680})$	[10]
23	NPCI	$(R_{680} - R_{430}) / (R_{680} + R_{430})$	[5]
24	PRI	$(R_{531} - R_{570}) / (R_{531} + R_{570})$	[11]
25	PRI_norm	$PRI * (-1) / (RDVI * R_{700} / R_{670})$	[12]
26	PRI*CI2	$PRI * CI2$	[13]
27	PSSR	R_{800} / R_{635}	[14]
28	PSND	$(R_{800} - R_{470}) / (R_{800} + R_{470})$	[14]
29	REP_LE	Red-edge position through linear extrapolation	[15]
30	SR7	R_{440} / R_{690}	[16]
31	SR8	R_{515} / R_{550}	[17]
32	SRWI	R_{850} / R_{1240}	[2]
33	Sum_Dr1	$\sum_{i=626}^{795} D1_i$	[18]
34	Sum_Dr2	$\sum_{i=680}^{780} D1_i$	[19]
35	Vogelmann	R_{740} / R_{720}	[20]
36	Vogelmann2	$(R_{734} - R_{747}) / (R_{715} + R_{726})$	[20]
37	D715/705	D_{715} / D_{705}	[20]
38	Vogelmann4	$(R_{734} - R_{747}) / (R_{715} + R_{720})$	[20]

Table S2. Detailed information on leaf samples.

Leaf Samples No.	Cultivar	Plant ID Per Cultivar	Type
1	8H534	2	healthy
2	8H534	3	healthy
3	DH351	1	healthy
4	DH351	3	healthy
5	Chang7-2	2	healthy

6	Chang7-2	3	healthy
7	8H953	5	healthy
8	8H953	6	healthy
9	PH6WC	7	healthy
10	PH6WC	8	healthy
11	DH351	5	healthy
12	DH351	6	healthy
13	Jing92	4	healthy
14	Jing92	8	healthy
15	Jing92	9	healthy
16	Jing724	5	healthy
17	Jing724	7	healthy
18	Jing724	8	healthy
19	8H534	6	Light
20	8H534	7	Light
21	DH351	7	Light
22	DH351	6	Light
23	Jing724	7	Light
24	Jing724	6	Light
25	Jing92	2	Light
26	Jing92	3	Light
27	DH382	3	Light
28	DH382	4	Light
29	Jing92	5	Light
30	Jing92	4	Light
31	Chang7-2	7	Light
32	Chang7-2	8	Light
33	PH6WC	4	Light
34	PH6WC	5	Light
35	8H534	8	Medium
36	8H953	9	Medium
37	Jing92	1	Medium
38	Jing92	4	Medium
39	Jing724	5	Medium
40	PH6WC	9	Medium
41	DH351	5	Medium
42	DH382	4	Medium
43	DH382	9	Medium
44	Chang7-2	7	Medium
45	Chang7-2	6	Medium
46	8H534	5	Severe
47	DH382	1	Severe
48	DH351	10	Severe
49	8H953	10	Severe
50	Chang7-2	6	Severe
51	Chang7-2	10	Severe
52	Jing724	3	Severe
53	Jing92	4	Severe
54	PH6WC	10	Severe

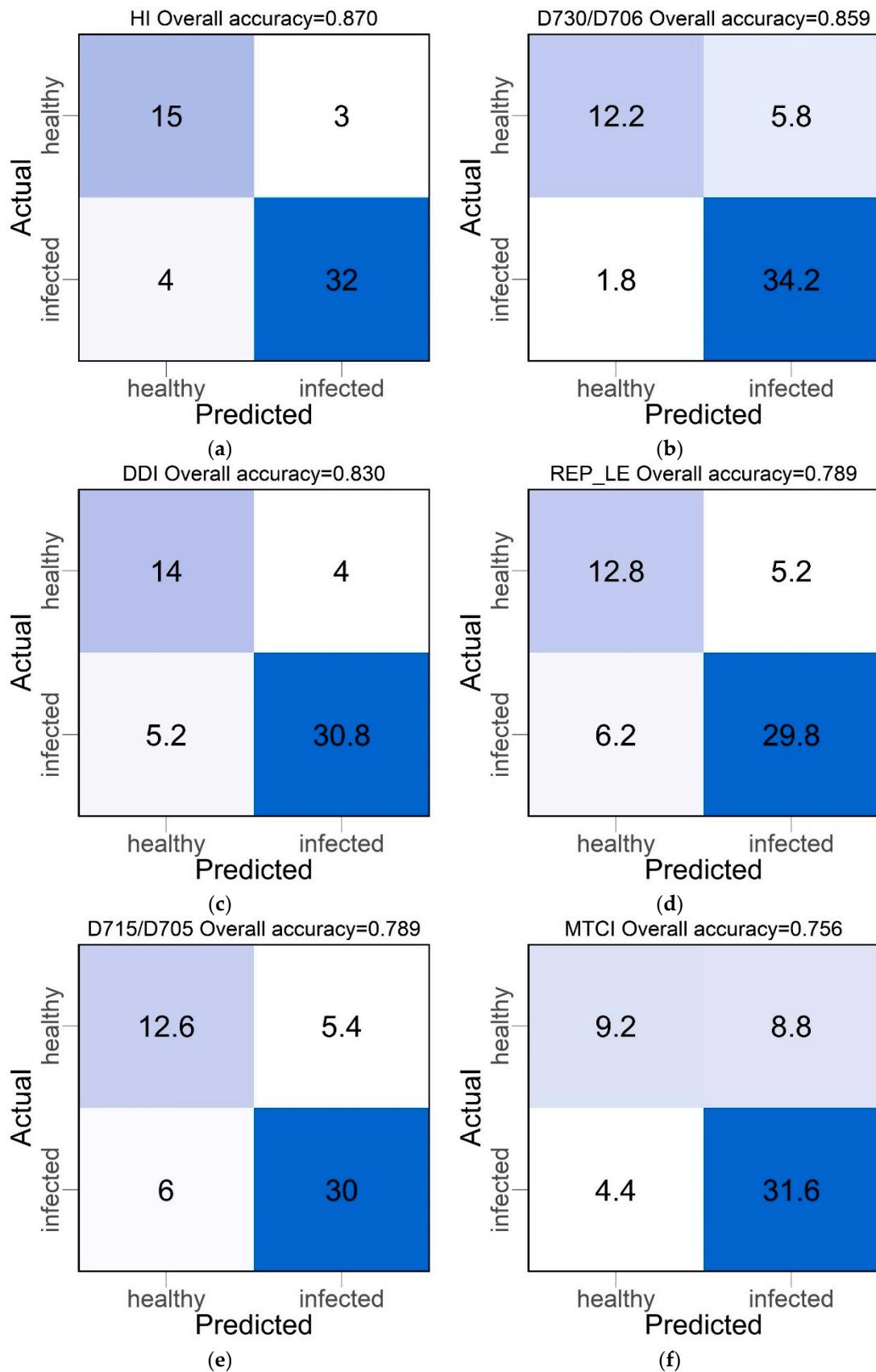


Figure S1. Averaged confusion matrix of different SCR detection models using support vector machine based 10-fold cross validation: a. Health Index (HI); b. The ratio of first derivative values at 730-706nm (D730/D706); c. Double Difference Index (DDI); d. The Red-edge position through linear extrapolation (REP_LE); e. The ratio of first derivative values at 715-705nm (D715/D705); f. MERIS

Terrestrial Chlorophyll Index (MTCI); the fraction number in the matrix was from the 10-fold cross validation results divided by ten.

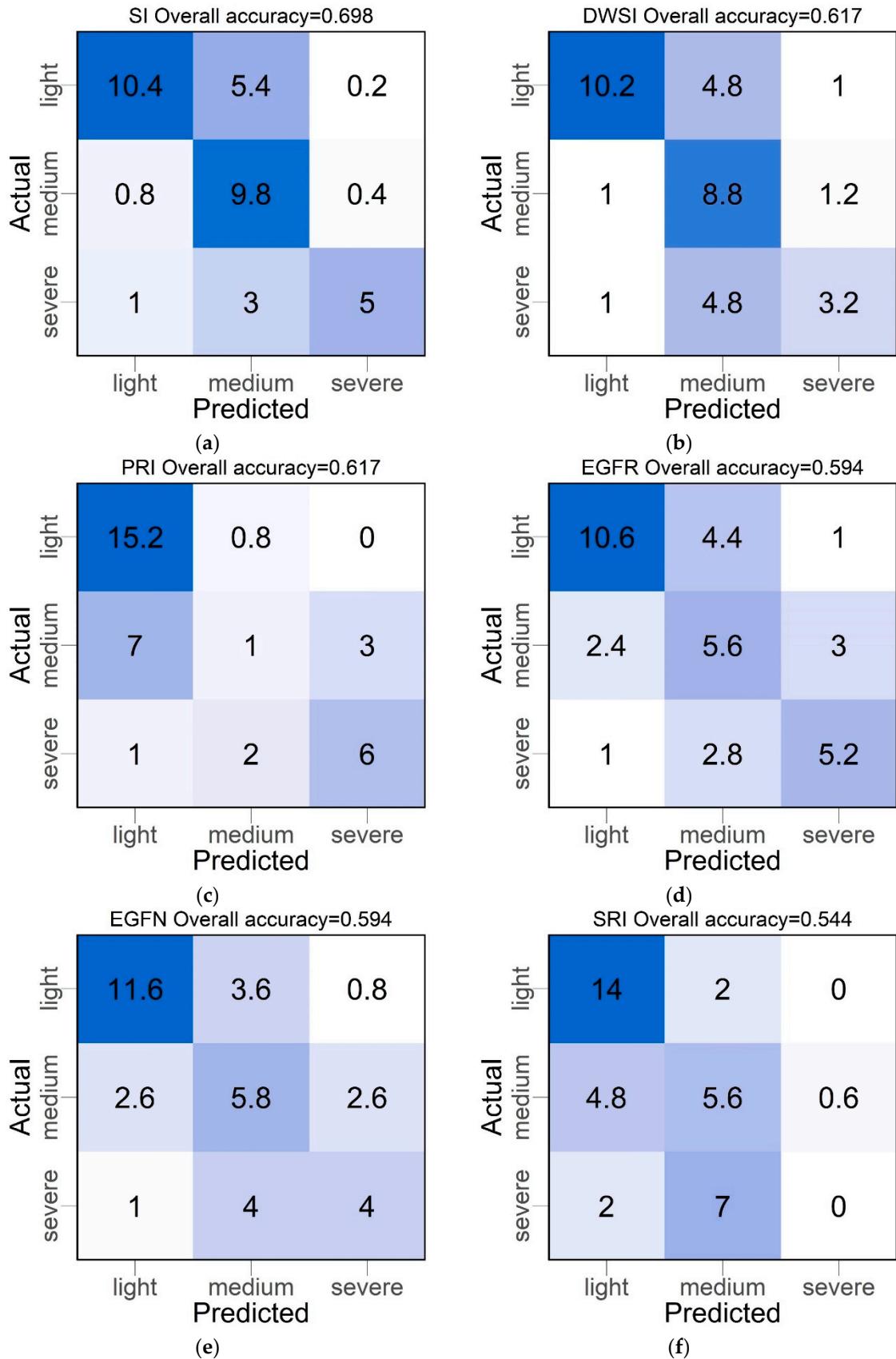


Figure S2. Averaged confusion matrix of different SCR severity models using support vector machine based 10-fold cross validation: a. Severity Index (SI); b. Disease-Water Stress Index (DWSI); c.

Physiological Reflectance Index (PRI); d. The simple ratio between the maxima of the first derivatives of reflectance at the red edge and green regions (EGFR); e. The normalized ratio between the maxima of the first derivatives of reflectance at the red edge and green regions (EGFN); f. Simple Ratio Index (SRI); the fraction number in the matrix was from the 10-fold cross validation results divided by ten.

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