## Supplementary tables and figures

## **Grapevine Phyllosphere Community Analysis in Response to Elicitor Application against Powdery Mildew**

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Supp	lementary	Table	<b>S1.</b> ]	Resume of	f read	counts and	l statistics	for each	sequenced	samp	ole in the	MiSeq	run
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Sample	Total Bases	Read Counts	GC (%)	AT (%)	Q20 (%)	Q30 (%)
CTRL1_MO_DNA_ITS	50,897,896	169,096	65.08	34.92	84.02	73.12
CTRL2_MO_DNA_ITS	50,721,510	168,510	65.30	34.70	83.70	72.87
CTRL3_MO_DNA_ITS	47,937,260	159,260	63.79	36.21	84.43	73.83
AcS-Mt1_MO_DNA_ITS	49,229,754	163,554	67.35	32.65	83.16	72.08
AcS-Mt2_MO_DNA_ITS	52,181,360	173,360	67.88	32.12	82.69	71.34
AcS-Mt3_MO_DNA_ITS	49,565,670	164,670	66.73	33.27	83.30	72.28
K-Pho1_MO_DNA_ITS	64,607,242	214,642	68.11	31.89	83.29	72.36
K-Pho2_MO_DNA_ITS	61,665,870	204,870	67.27	32.73	83.56	72.52
K-Pho3_MO_DNA_ITS	60,049,500	199 <i>,</i> 500	67.33	32.67	83.93	73.04
Lam1_MO_DNA_ITS	65,298,940	216,940	65.61	34.39	84.27	73.50
Lam2_MO_DNA_ITS	50,214,626	166,826	67.32	32.68	83.41	72.28
Lam3_MO_DNA_ITS	50,790,740	168,740	68.46	31.54	83.52	72.67
CTRL1_NE_DNA_ITS	50,250,144	166,944	63.98	36.02	83.28	72.33
CTRL2_NE_DNA_ITS	61,148,752	203,152	63.38	36.62	84.50	73.57
CTRL3_NE_DNA_ITS	58,818,410	195,410	64.46	35.54	83.29	72.17
AcS-Mt1_NE_DNA_ITS	55,802,992	185,392	67.0	33.0	82.60	71.31
AcS-Mt2_NE_DNA_ITS	51,794,876	172,076	66.12	33.88	83.0	71.78
AcS-Mt3_NE_DNA_ITS	56,359,842	187,242	66.57	33.43	82.90	71.80
K-Pho1_NE_DNA_ITS	60,026,624	199,424	65.96	34.04	81.36	69.48
K-Pho2_NE_DNA_ITS	63,471,870	210,870	64.90	35.10	82.88	71.92
K-Pho3_NE_DNA_ITS	53,839,268	178,868	67.06	32.94	82.61	71.44
Lam1_NE_DNA_ITS	57,239,966	190,166	65.36	34.64	84.31	73.44
Lam2_NE_DNA_ITS	54,599,594	181,394	64.01	35.99	83.99	73.06
Lam3_NE_DNA_ITS	51,449,328	170,928	65.75	34.25	82.74	71.41

	NE CTRL	NE AcS-Mt	NE K-Pho	NE Lam	MO CTRL	MO AcS-Mt	MO K-Pho	MO Lam
Genera	Mean Sl	D Mean SD	Mean SD	Mean SD	Mean SD	Mean SD	Mean SD	Mean SD
Erysiphe	77.71% ± 6.86	% 74.73% ± 5.31%	67.44% ± 10.29%	86.71% ± 2.11%	$50.78\% \pm 10.45\%$	$22.28\% \pm 6.18\%$	$21.52\% \pm 6.61\%$	$38.90\% \pm 5.97\%$
Alternaria	$10.08\% \pm 2.88$	% 10.39% ± 4.43%	13.48% ± 3.28%	$1.02\% \pm 0.55\%$	$20.72\% \pm 2.81\%$	35.33% ± 3.58%	32.87% ± 3.88%	$22.52\% \pm 1.90\%$
Cladosporium	$5.23\% \pm 0.83$	$\% 5.10\% \pm 0.04\%$	7.09% ± 2.58%	$3.75\% \pm 1.14\%$	$10.31\% \pm 4.63\%$	$12.40\% \pm 3.72\%$	$12.98\% \pm 2.06\%$	$16.43\% \pm 2.13\%$
Epicoccum	$5.27\% \pm 1.92$	$\% 6.55\% \pm 0.66\%$	$8.56\% \pm 1.14\%$	2.48% ± 1.33%	$12.19\% \pm 1.21\%$	13.28% ± 3.09%	$24.58\% \pm 3.90\%$	$10.48\% \pm 1.13\%$
Aureobasidium	$0.26\% \pm 0.10$	$\% 0.22\% \pm 0.13\%$	$0.72\% \pm 0.20\%$	$0.16\% \pm 0.06\%$	$0.92\% \pm 0.51\%$	$10.01\% \pm 1.79\%$	$2.25\% \pm 1.10\%$	$2.16\% \pm 0.74\%$
Bipolaris	$0.56\% \pm 0.29$	$\% 0.53\% \pm 0.44\%$	$0.43\% \pm 0.19\%$	$0.46\% \pm 0.30\%$	$1.69\% \pm 0.55\%$	$2.32\% \pm 0.43\%$	$1.86\% \pm 0.26\%$	$5.44\% \pm 0.29\%$
Pithomyces	$0.32\% \pm 0.12$	% 1.52% ± 0.26%	$1.29\% \pm 0.28\%$	$0.85\% \pm 0.40\%$	$1.86\% \pm 0.76\%$	$2.44\% \pm 1.44\%$	$1.91\% \pm 0.46\%$	$0.96\% \pm 0.38\%$
Filobasidium	$0.12\% \pm 0.02$	$\% 0.02\% \pm 0.01\%$	$0.20\% \pm 0.04\%$	$0.03\% \pm 0.01\%$	$0.89\% \pm 0.52\%$	$0.25\% \pm 0.13\%$	$0.20\% \pm 0.08\%$	$0.07\% \pm 0.03\%$
Curvularia	$0.22\% \pm 0.12$	$\% 0.26\% \pm 0.17\%$	$0.28\% \pm 0.06\%$	$0.11\% \pm 0.09\%$	$0.15\% \pm 0.10\%$	$0.50\% \pm 0.06\%$	$0.27\% \pm 0.11\%$	$0.91\% \pm 0.10\%$
Nigrospora	$0.22\% \pm 0.19$	$\% 0.68\% \pm 0.07\%$	$0.50\% \pm 0.29\%$	$0.43\% \pm 0.16\%$	$0.49\% \pm 0.11\%$	$1.20\% \pm 0.36\%$	$1.55\% \pm 0.96\%$	$2.12\% \pm 0.74\%$

**Supplementary Table S2.** Abundances of TOP 10 fungal genera among the analyzed samples

**Supplementary Table S3.** List of OTUs identified in 'Moscato' (MO) samples. Values are the mean of three biological replicates ± standard deviation (SD). MO CTRL = non-treated inoculated plants; MO AcS-Mt = plants treated with Acibenzolar-S-methyl (Bion, Syngenta Crop Protection); MO K-Pho = plants treated with Potassium phosphonate (Century, BASF Agro); MO Lam = inoculated plants treated with Laminarin (Vacciplant, Arysta Lifescience).

	MOC	TRL	MO A	cS-Mt	MO K	-Pho	МО	) Lam	
	Mean	SD	Mean	SD	Mean	SD	Mean	SD	
Erysiphe	49.96% ±	30.29%	21.84% ±	= 17.63%	19.84% ±	17.82%	38.47% :	± 27.95%	
Epicoccum	11.87% ±	9.83%	13.09% ±	3.19%	22.37% ±	3.78%	10.34% :	± 7.16%	
Alternaria	20.26% ±	12.48%	34.75% ±	: 13.19%	29.84% ±	12.84%	22.08% =	± 7.27%	
Cladosporium	10.08% ±	4.45%	12.18% ±	3.52%	11.76% ±	1.39%	16.05% :	± 11.58%	
Aureobasidium	0.90% ±	0.50%	9.99% ±	: 15.78%	2.06% ±	1.07%	2.16% :	± 3.74%	
Nigrospora	0.48% ±	0.41%	1.18% ±	0.36%	1.41% ±	0.90%	2.07% :	± 1.95%	
Claviceps	0.00% ±	0.00%	0.00% ±	- 0.00%	0.57% ±	0.95%	0.00% :	± 0.00%	
Phoma	0.00% ±	0.00%	0.00% ±	- 0.00%	0.62% ±	0.90%	0.00% :	± 0.00%	
Bipolaris	1.65% ±	1.50%	2.28% ±	- 0.45%	1.70% ±	0.24%	5.42% :	± 8.29%	
Coniozyma	0.00% ±	0.00%	0.00% ±	- 0.00%	0.32% ±	0.56%	0.00% :	± 0.00%	
Hannaella	0.12% ±	0.22%	0.14% ±	- 0.24%	0.55% ±	0.34%	0.00% :	± 0.00%	
Curvularia	0.15% ±	0.20%	0.49% ±	- 0.66%	0.24% ±	0.26%	0.89% :	± 0.79%	
Ulocladium	0.00% ±	0.00%	0.00% ±	- 0.00%	0.23% ±	0.23%	0.90% :	± 1.51%	
Pithomyces	1.82% ±	0.73%	2.39% ±	= 1.40%	1.69% ±	1.22%	0.93% :	± 1.33%	
Fusarium	0.00% ±	0.00%	0.41% ±	- 0.49%	0.39% ±	0.53%	0.06% :	± 0.05%	
Lophiostoma	0.00% ±	0.00%	0.00% ±	- 0.00%	0.24% ±	0.37%	0.00% :	± 0.00%	
Blumeria	0.00% ±	0.00%	0.00% ±	- 0.00%	0.00% ±	0.00%	0.12% :	± 0.10%	
Candida	0.00% ±	0.00%	0.12% ±	- 0.20%	0.00% ±	0.00%	0.00% :	± 0.00%	
Filobasidium	0.87% ±	1.01%	0.24% ±	- 0.42%	0.17% ±	0.23%	0.00% :	± 0.00%	
Leptosphaerulina	0.44% ±	0.75%	0.00% ±	- 0.00%	0.00% ±	0.00%	0.00% :	± 0.00%	
Microsphaeropsis	0.00% ±	0.00%	0.30% ±	- 0.37%	0.76% ±	0.82%	0.00% :	± 0.00%	
Noosia	0.00% ±	0.00%	0.00% ±	- 0.00%	0.14% ±	0.12%	0.00% :	± 0.00%	
Periconia	0.34% ±	0.47%	0.12% ±	- 0.21%	0.00% ±	0.00%	0.00% :	± 0.00%	
Pleochaeta	0.30% ±	0.07%	0.00% ±	- 0.00%	0.00% ±	0.00%	0.00% :	± 0.00%	
Podospora	0.00% ±	0.00%	0.00% ±	- 0.00%	0.00% ±	0.00%	0.45% :	± 0.54%	
Preussia	0.42% ±	0.44%	0.34% ±	- 0.42%	0.22% ±	0.39%	0.00% :	± 0.00%	
Stemphylium	0.15% ±	0.13%	0.00% ±	- 0.00%	4.51% ±	7.80%	0.00% :	± 0.00%	
Symmetrospora	0.17% ±	0.26%	0.00% ±	- 0.00%	0.00% ±	0.00%	0.00% :	± 0.00%	
Trichoderma	0.00% ±	0.00%	0.15% ±	- 0.11%	0.00% ±	0.00%	0.05% :	± 0.09%	
Vishniacozyma	0.00% ±	0.00%	0.00% ±	= 0.00%	0.36% ±	0.44%	0.00% :	± 0.00%	

**Supplementary Table S4.** List of OTUs identified in 'Nebbiolo' (NE) samples. Values are the mean of three biological replicates ± standard deviation (SD). CTRL = non-treated inoculated plants; AcS-Mt = plants treated with Acibenzolar-S-methyl (Bion, Syngenta Crop Protection); K-Pho = plants treated with Potassium phosphonate (Century, BASF Agro); Lam = inoculated plants treated with Laminarin (Vacciplant, Arysta Lifescience).

	NE CTRL	NE AcS-Mt	NE K-Pho	NE Lam
	Mean SD	Mean SD	Mean SD	Mean SD
Erysiphe	$68.76\% \pm 24.88\%$	$30.32\% \pm 1.98\%$	$36.37\% \pm 20.44\%$	63.68% ± 18.73%
Alternaria	$7.76\% \pm 1.84\%$	$4.39\% \pm 2.28\%$	$5.92\% \pm 1.97\%$	$3.60\% \pm 0.57\%$
Cladosporium	$4.58\% \pm 1.73\%$	$2.08\% \pm 0.26\%$	$3.44\% \pm 1.52\%$	$2.67\% \pm 0.80\%$
Epicoccum	$4.30\% \pm 0.99\%$	$2.68\% \pm 0.53\%$	$4.26\% \pm 1.42\%$	$1.69\% \pm 0.56\%$
Aureobasidium	$0.23\% \pm 0.17\%$	$0.09\% \pm 0.06\%$	$0.48\% \pm 0.69\%$	$0.11\% \pm 0.04\%$
Bipolaris	$0.51\% \pm 0.37\%$	$0.23\% \pm 0.21\%$	$0.22\% \pm 0.23\%$	$0.38\% \pm 0.31\%$
Pithomyces	$0.31\% \pm 0.25\%$	$0.63\% \pm 0.17\%$	$0.69\% \pm 0.34\%$	$0.58\% \pm 0.15\%$
Filobasidium	$0.12\% \pm 0.17\%$	$0.01\% \pm 0.01\%$	$0.14\% \pm 0.17\%$	$0.02\% \pm 0.01\%$
Curvularia	$0.22\% \pm 0.24\%$	$0.10\% \pm 0.05\%$	$0.09\% \pm 0.10\%$	$0.08\% \pm 0.08\%$
Nigrospora	$0.21\% \pm 0.21\%$	$0.28\% \pm 0.03\%$	$0.19\% \pm 0.16\%$	$0.31\% \pm 0.11\%$
Pleochaeta	$0.15\% \pm 0.17\%$	$0.05\% \pm 0.05\%$	$0.13\% \pm 0.12\%$	$0.13\% \pm 0.10\%$
Ulocladium	$0.14\% \pm 0.06\%$	$0.06\% \pm 0.06\%$	$0.01\% \pm 0.02\%$	$0.09\% \pm 0.11\%$
Blumeria	$0.08\% \pm 0.01\%$	$0.03\% \pm 0.01\%$	$0.04\% \pm 0.03\%$	$0.08\% \pm 0.05\%$
Hannaella	$0.11\% \pm 0.07\%$	$0.00\% \pm 0.01\%$	$0.04\% \pm 0.04\%$	$0.03\% \pm 0.03\%$
Leptosphaerulina	$0.04\% \pm 0.03\%$	$0.03\% \pm 0.03\%$	$0.02\% \pm 0.01\%$	$0.04\% \pm 0.05\%$
Vishniacozyma	$0.02\% \pm 0.03\%$	$0.00\% \pm 0.01\%$	$0.18\% \pm 0.29\%$	$0.01\% \pm 0.02\%$
Fusarium	$0.06\% \pm 0.06\%$	$0.07\% \pm 0.05\%$	$0.44\% \pm 0.64\%$	$0.14\% \pm 0.12\%$
Periconia	$0.02\% \pm 0.02\%$	$0.09\% \pm 0.10\%$	$0.15\% \pm 0.20\%$	$0.03\% \pm 0.02\%$
Microsphaeropsis	$0.04\% \pm 0.04\%$	$0.05\% \pm 0.05\%$	$0.19\% \pm 0.18\%$	$0.00\% \pm 0.00\%$
Stemphylium	$0.03\% \pm 0.03\%$	$0.01\% \pm 0.03\%$	$0.02\% \pm 0.02\%$	$0.05\% \pm 0.05\%$
Sordaria	$0.00\% \pm 0.00\%$	$0.10\% \pm 0.17\%$	$0.00\% \pm 0.00\%$	$0.00\% \pm 0.00\%$

**Supplementary Table S5.** Bray-Curtis matrix was used to perform PERMANOVA analysis to highlight the genotype effect. All samples from each genotype were considered independently of treatment type. *p* (Bonferroni-corrected) and *F* values for comparison of 'Nebbiolo' (NE) and 'Moscato' (MO) samples are reported below the table.

	NE	МО
<i>P</i> value		
NE		0.0001
MO	0.0001	-
F: 23.16		-

**Supplementary Table S6.** Bray-Curtis matrix was used to perform PERMANOVA analysis to assess possible effects of treatment on 'Nebbiolo' (NE). *p* (Bonferroni-corrected) and *F* values are reported below the table. CTRL = non-treated inoculated plants; AcS-Mt = plants treated with Acibenzolar-S-methyl (Bion, Syngenta Crop Protection); K-Pho = plants treated with Potassium phosphonate (Century, BASF Agro); Lam = inoculated plants treated with Laminarin (Vacciplant, Arysta Lifescience).

	NE CTRL	NE AcS-Mt	NE K-Pho	NE Lam
P value				
NE CTRL		0.499	0.2074	0.1042
NE AcS-Mt	0.499		0.3988	0.1055
NE K-Pho	0.2074	0.3988		0.101
NE Lam	0.1042	0.1055	0.101	
F value				
NE CTRL		0.5514	1.526	3.382
NE Ac-Mt	0.5514		0.8985	9.57
NE K-Pho	1.526	0.8985		7.114
NE Lam	3.382	9.57	7.114	

**Supplementary Table S7.** Bray-Curtis matrix was used to perform PERMANOVA analysis to assess possible effects of treatment on 'Moscato' (MO). *p* (Bonferroni-corrected) and *F* values are reported below the table. CTRL = non-treated inoculated plants; AcS-Mt = plants treated with Acibenzolar-S-methyl (Bion, Syngenta Crop Protection); K-Pho = plants treated with Potassium phosphonate (Century, BASF Agro); Lam = inoculated plants treated with Laminarin (Vacciplant, Arysta Lifescience).

	MO CTRL	MO AcS-Mt	MO K-Pho	MO Lam
<i>P</i> value				
MO CTRL		0.2999	0.3025	0.6088
MO AcS-Mt	0.2999		0.7008	0.6097
MO K-Pho	0.3025	0.7008		0.4067
MO Lam	0.6088	0.6097	0.4067	
F value				
MO CTRL		1.67	1.874	0.2504
MO Ac-Mt	1.67		0.4393	0.8533
MO K-Pho	1.874	0.4393		1.055
MO Lam	0.2504	0.8533	1.055	

**Supplementary Table S8.** Bray-Curtis matrix was used to perform PERMANOVA analysis in order to compare treatment effects on 'Nebbiolo' (NE) and 'Moscato' (MO) samples at once. *p* (Bonferroni-corrected) and *F* values are reported below the table. CTRL = non-treated inoculated plants; AcS-Mt = plants treated with Acibenzolar-S-methyl (Bion, Syngenta Crop Protection); K-Pho = plants treated with Potassium phosphonate (Century, BASF Agro); Lam = inoculated plants treated with Laminarin (Vacciplant, Arysta Lifescience).

	NE CTRL	NE AcS-Mt	NE K-Pho	NE Lam	MO CTRL	MO AcS-Mt	MO K-Pho	MO Lam
P value								
NE CTRL		0.0988	0.1941	0.7985	0.0946	0.1058	0.1019	0.2002
NE AcS-Mt	0.0988		0.7074	0.1035	0.405	0.0978	0.0989	0.4068
NE K-Pho	0.1941	0.7074		0.1998	0.3083	0.102	0.0988	0.2991
NE Lam	0.7985	0.1035	0.1998		0.1002	0.0999	0.0984	0.1967
MO CTRL	0.0946	0.405	0.3083	0.1002		0.3067	0.2997	1
MO AcS-Mt	0.1058	0.0978	0.102	0.0999	0.3067		0.7905	0.9029
MO K-Pho	0.1019	0.0989	0.0988	0.0984	0.2997	0.7905		0.7969
MO Lam	0.2002	0.4068	0.2991	0.1967	1	0.9029	0.7969	
F value								
NE CTRL		8.552	1.672	0.8406	3.89	10.54	10.45	3.399
NE Ac-Mt	8.552		0.6059	10.94	1.981	9.368	10.54	2.335
NE K-Pho	1.672	0.6059		2.146	1.083	4.463	4.413	1.72
NE Lam	0.8406	10.94	2.146		4.373	13.6	15.03	3.408
MO CTRL	3.89	1.981	1.083	4.373		1.054	1.215	0.3847
MO Ac-Mt	10.54	9.368	4.463	13.6	1.054		0.4001	0.5917
MO K-Pho	10.45	10.54	4.413	15.03	1.215	0.4001		0.7494
MO Lam	3.399	2.335	1.72	3.408	0.3847	0.5917	0.7494	

**Supplementary Table S9.** Alpha diversity indices calculated starting from ITS sequencing data obtained for each treatment in 'Nebbiolo' (NE) and 'Moscato' (MO). CTRL = non-treated inoculated plants; AcS-Mt = plants treated with Acibenzolar-S-methyl (Bion, Syngenta Crop Protection); K-Pho = plants treated with Potassium phosphonate (Century, BASF Agro); Lam = inoculated plants treated with Laminarin (Vacciplant, Arysta Lifescience).

_	NE CTI	RL	NE AcS-	·Mt	NE K-P	ho	NE Lai	m	MO CT	RL	MO AcS	-Mt	MO K-I	Pho	MO La	m
	Mean	SD	Mean	SD	Mean	SD	Mean	SD	Mean	SD	Mean	SD	Mean	SD	Mean	SD
Simpson (D)	0.38 ±	0.09	$0.44 \pm$	0.06	0.53 ±	0.11	0.26 ±	0.04	0.60 ±	0.28	0.74 ±	0.05	0.76 ±	0.05	0.69 ±	0.19
Shannon (H')	0.87 ±	0.13	1.01 ±	0.10	1.18 ±	0.16	$0.65 \pm$	0.09	1.37 ±	0.60	1.64 ±	0.10	1.72 ±	0.17	1.49 ±	0.45
Chao	20.33 ±	1.53	20.33 ±	1.00	$20.67 \pm$	3.21	19.67 ±	1.00	20.33 ±	3.61	19.67 ±	2.31	19.67 ±	2.08	$20.00 \pm$	2.00

1-         1 $1_{MO}_{RNA}$ $51,442,584$ $46.75$ $53.25$ $98.74$ $95.99$ $1_{NE}_{RNA}$ $49,070,822$ $46.68$ $53.32$ $98.75$ $95.97$ $1_{-}$ $2_{MO}_{RNA}$ $45,950,076$ $46.20$ $53.80$ $98.68$ $95.78$ $1_{-}$ $3_{-}$ $52,359,574$ $46.09$ $53.91$ $98.73$ $95.96$ $1_{-}$ $3_{-}$ $52,359,574$ $46.09$ $53.91$ $98.73$ $95.96$ $1_{-}$ $3_{-}$ $52,359,574$ $46.40$ $53.60$ $98.75$ $96.0$ $2_{-}$ $2_{-}$ $2_{-}$ $46.46$ $53.54$ $98.62$ $95.62$ $2_{-}$ $2_{-}$ $2_{-}$ $2_{-}$ $2_{-}$ $2_{-}$ $2_{-}$ $2_{-}$ $2_{-}$ $2_{-}$ $2_{-}$ $2_{-}$ $2_{-}$ $2_{-}$ $2_{-}$ $2_{-}$ $2_{-}$ $2_{-}$ $2_{-}$ $45.87$ $54.13$ $98.79$ $95.82$ $2_{-}$ <th>Sample</th> <th>Read Counts</th> <th>GC (%)</th> <th>AT (%)</th> <th>Q20 (%)</th> <th>Q30 (%)</th>	Sample	Read Counts	GC (%)	AT (%)	Q20 (%)	Q30 (%)
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	1-					
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	1 MO RNA	51,442,584	46.75	53.25	98.74	95.99
1- $2\_MO\_RNA$ 45,950,076       46.20       53.80       98.68       95.78 $1-2\_NE\_RNA$ 52,359,574       46.09       53.91       98.73       95.96         1-       3_MO\_RNA       44,637,026       46.27       53.74       98.72       95.93 $1-3\_NE\_RNA$ 105,207,854       46.46       53.54       98.62       95.62 $2-$ -       -       -       -       98.75       96.0 $2-1\_NE\_RNA$ 90,822,934       45.94       54.06       98.70       95.82 $2$ -       -       -       -       - $2\_MO\_RNA$ 45,539,430       48.83       51.17       98.76       95.82 $2$ -       -       -       -       -       - $3\_MO\_RNA$ 50,716,752       46.26       53.74       98.66       95.85       - $2$ -       -	1-1 NE RNA	49,070,822	46.68	53.32	98.75	95.97
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	1-	, ,				
1-2_NE_RNA 52,359,574 46.09 53.91 98.73 95.96 1- 3_MO_RNA 44,637,026 46.27 53.74 98.72 95.93 1-3_NE_RNA 105,207,854 46.46 53.54 98.62 95.62 2- 1_MO_RNA 43,016,122 46.40 53.60 98.75 96.0 2-1_NE_RNA 99,822,934 45.94 54.06 98.70 95.82 2- 2_MO_RNA 45,539,430 48.83 51.17 98.71 95.88 2-2_NE_RNA 98,299,580 45.87 54.13 98.76 95.93 2- 3_MO_RNA 50,716,752 46.26 53.74 98.66 95.85 2-3_NE_RNA 98,607,242 45.75 54.26 98.69 95.75 3- 1_MO_RNA 49,042,662 46.46 53.54 98.67 95.82 3NE_RNA 82,768,782 45.87 54.13 98.59 95.61 3- 2_MO_RNA 51,651,078 46.35 53.65 98.85 96.31 3-2_NE_RNA 87,883,904 46.60 53.40 98.59 95.59 3- 3_MO_RNA 54,955,000 46.32 53.68 98.84 96.34 3-3_NE_RNA 52,636,722 47.58 52.42 98.66 95.84 4- 1_MO_RNA 46,153,778 46.05 53.95 98.86 96.30 4-1_NE_RNA 53,996,182 46.73 53.27 98.83 96.33 4- 2_MO_RNA 42,770,800 46.44 53.57 98.83 96.33 4-2_NO_RNA 42,770,800 46.44 53.57 98.83 96.33 4-2_MO_RNA 42,770,800 46.44 53.57 98.84 96.34 5-2_NE_RNA 48,834,504 46.25 53.75 98.78 96.07 5- 1_MO_RNA 94,239,798 46.40 53.60 98.68 95.97 5-1_NE_RNA 48,834,504 46.25 53.75 98.78 96.07 5- 2_MO_RNA 99,231,268 46.44 53.56 98.74 96.03 5-2_NE_RNA 41.094,700 45.74 54 26 98.75 95.93	2 MO RNA	45,950,076	46.20	53.80	98.68	95.78
1-         1-           3_MO_RNA         44,637,026         46.27         53.74         98.72         95.93           1-3_NE_RNA         105,207,854         46.46         53.54         98.62         95.62           2-         1_MO_RNA         43,016,122         46.40         53.60         98.75         96.0           2-1_NE_RNA         99,822,934         45.94         54.06         98.70         95.82           2-         2         -         -         -         -           2_MO_RNA         45,539,430         48.83         51.17         98.71         95.88           2-2         -         -         -         -         -         -           3_MO_RNA         50,716,752         46.26         53.74         98.66         95.85           2-3_NE_RNA         98,607,242         45.75         54.26         98.67         95.82           3_MO_RNA         50,716,752         46.26         53.54         98.67         95.82           3_NE_RNA         98,607,242         45.87         54.13         98.59         95.61           3_         -         -         -         -         -         -           2_MO_RNA <td>1-2 NE RNA</td> <td>52.359.574</td> <td>46.09</td> <td>53.91</td> <td>98.73</td> <td>95.96</td>	1-2 NE RNA	52.359.574	46.09	53.91	98.73	95.96
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	1-	- , ,-				
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	3 MO RNA	44.637.026	46.27	53.74	98.72	95.93
2- 1_MO_RNA 43,016,122 46.40 53.60 98.75 96.0 2-1_NE_RNA 99,822,934 45.94 54.06 98.70 95.82 2- 2_MO_RNA 45,539,430 48.83 51.17 98.71 95.88 2-2_NE_RNA 98,299,580 45.87 54.13 98.76 95.93 2- 3_MO_RNA 50,716,752 46.26 53.74 98.66 95.85 2-3_NE_RNA 98,607,242 45.75 54.26 98.69 95.75 3- 1_MO_RNA 49,042,662 46.46 53.54 98.67 95.82 3-1_NE_RNA 82,768,782 45.87 54.13 98.59 95.61 3- 2_MO_RNA 51,651,078 46.35 53.65 98.85 96.31 3-2_NE_RNA 82,768,722 47.58 52.42 98.66 95.84 4- 1_MO_RNA 54,955,000 46.32 53.68 98.84 96.34 3-3_NE_RNA 52,636,722 47.58 52.42 98.66 95.84 4- 1_MO_RNA 46,153,778 46.05 53.95 98.86 96.30 4-1_NE_RNA 53,996,182 46.73 53.27 98.67 95.93 4- 2_MO_RNA 42,770,800 46.44 53.57 98.83 96.33 4-2_NE_RNA 53,019,634 46.91 53.09 98.69 95.81 4- 3_MO_RNA 109,690,716 46.39 53.61 98.85 96.19 4- 3_MO_RNA 47,729,898 46.34 53.66 98.71 95.92 5- 1_MO_RNA 94,239,798 46.40 53.60 98.68 95.97 5- 1_MO_RNA 94,239,798 46.40 53.60 98.68 95.97 5- 1_MO_RNA 48,834,504 46.25 53.75 98.78 96.07 5- 2_MO_RNA 99,231,268 46.44 53.56 98.74 96.03 5-2 NE_RNA 41,094,770 45.74 54.26 98.75 95.92	1-3 NE RNA	105,207,854	46.46	53.54	98.62	95.62
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	2-	, ,				
2-1_NE_RNA 99,822,934 45.94 54.06 98.70 95.82 2- 2_MO_RNA 45,539,430 48.83 51.17 98.71 95.88 2-2_NE_RNA 98,299,580 45.87 54.13 98.76 95.93 2- 3_MO_RNA 50,716,752 46.26 53.74 98.66 95.85 2-3_NE_RNA 98,607,242 45.75 54.26 98.69 95.75 3- 1_MO_RNA 49,042,662 46.46 53.54 98.67 95.82 3-1_NE_RNA 82,768,782 45.87 54.13 98.59 95.61 3- 2_MO_RNA 51,651,078 46.35 53.65 98.85 96.31 3-2_NE_RNA 87,883,904 46.60 53.40 98.59 95.59 3- 3_MO_RNA 54,955,000 46.32 53.68 98.84 96.34 3-3_NE_RNA 52,636,722 47.58 52.42 98.66 95.84 4- 1_MO_RNA 46,153,778 46.05 53.95 98.86 96.30 4-1_NE_RNA 53,996,182 46.73 53.27 98.67 95.93 4- 2_MO_RNA 42,770,800 46.44 53.57 98.83 96.33 4-2_NE_RNA 53,019,634 46.91 53.09 98.69 95.81 4- 3_MO_RNA 109,690,716 46.39 53.61 98.85 96.19 4- 3_MO_RNA 109,690,716 46.39 53.61 98.85 96.19 4- 3_MO_RNA 109,690,716 46.39 53.61 98.85 96.19 4- 3_MO_RNA 94,239,798 46.40 53.60 98.68 95.97 5- 1_MO_RNA 94,239,798 46.40 53.60 98.68 95.97 5- 2_MO_RNA 94,239,798 46.40 53.60 98.68 95.97 5- 2_MO_RNA 99,231,268 46.44 53.57 98.78 96.03 5-2 NE_RNA 41 094 770 45 74 54 26 98.74 96.03	1 MO RNA	43.016.122	46.40	53.60	98.75	96.0
2-       1	2-1 NE RNA	99.822.934	45.94	54.06	98.70	95.82
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	2-	, - , - ,				
2-2_NE_RNA 98,299,580 45.87 54.13 98.76 95.93 2- 3_MO_RNA 50,716,752 46.26 53.74 98.66 95.85 2-3_NE_RNA 98,607,242 45.75 54.26 98.69 95.75 3- 1_MO_RNA 49,042,662 46.46 53.54 98.67 95.82 3-1_NE_RNA 82,768,782 45.87 54.13 98.59 95.61 3- 2_MO_RNA 51,651,078 46.35 53.65 98.85 96.31 3-2_NE_RNA 87,883,904 46.60 53.40 98.59 95.59 3- 3_MO_RNA 54,955,000 46.32 53.68 98.84 96.34 3-3_NE_RNA 52,636,722 47.58 52.42 98.66 95.84 4- 1_MO_RNA 46,153,778 46.05 53.95 98.86 96.30 4-1_NE_RNA 53,996,182 46.73 53.27 98.87 95.93 4- 2_MO_RNA 42,770,800 46.44 53.57 98.83 96.33 4-2_NE_RNA 53,019,634 46.91 53.09 98.69 95.81 4- 3_MO_RNA 109,690,716 46.39 53.61 98.85 96.19 4-3_NE_RNA 109,690,716 46.39 53.61 98.85 96.19 4-3_NE_RNA 47,729,898 46.34 53.66 98.71 95.92 5- 1_MO_RNA 94,239,798 46.40 53.60 98.68 95.97 5-1_NE_RNA 48,834,504 46.25 53.75 98.78 96.03 5-2_NE_RNA 99,231,268 46.44 53.56 98.74 96.03 5-2_NE_RNA 41.094.770 45.74 54.26 98.75 95.92	2 MO RNA	45,539,430	48.83	51.17	98.71	95.88
2-       2-       2-         3_MO_RNA       50,716,752       46.26       53.74       98.66       95.85         2-3_NE_RNA       98,607,242       45.75       54.26       98.69       95.75         3-       1_MO_RNA       49,042,662       46.46       53.54       98.67       95.82         3-1_NE_RNA       82,768,782       45.87       54.13       98.59       95.61         3-       2_MO_RNA       51,651,078       46.35       53.65       98.85       96.31         3-2_NE_RNA       87,883,904       46.60       53.40       98.59       95.59         3-       3_MO_RNA       54,955,000       46.32       53.68       98.84       96.34         3-3_NE_RNA       52,636,722       47.58       52.42       98.66       95.84         4-       4-       4-       4-       4-       4-         1_MO_RNA       46,153,778       46.05       53.95       98.86       96.30         4-1_NE_RNA       53,019,634       46.91       53.09       98.69       95.81         4-       3_MO_RNA       109,690,716       46.39       53.61       98.85       96.19         4-3_NE_RNA       47,729,898       <	2-2 NE RNA	98,299.580	45.87	54.13	98.76	95.93
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	2-					
2-3_NE_RNA       98,607,242       45.75       54.26       98.69       95.75         3-       1_MO_RNA       49,042,662       46.46       53.54       98.67       95.82         3-1_NE_RNA       82,768,782       45.87       54.13       98.59       95.61         3-       2_MO_RNA       51,651,078       46.35       53.65       98.85       96.31         3-2_NE_RNA       87,883,904       46.60       53.40       98.59       95.59         3-       3_MO_RNA       54,955,000       46.32       53.68       98.84       96.34         3-3_NE_RNA       52,636,722       47.58       52.42       98.66       95.84         4-       1_MO_RNA       46,153,778       46.05       53.95       98.86       96.30         4-1_NE_RNA       53,996,182       46.73       53.27       98.67       95.93       4.         2_MO_RNA       42,770,800       46.44       53.57       98.83       96.33         4-2_NE_RNA       53,019,634       46.91       53.09       98.69       95.81         4-       -       -       -       -       -       -         3_MO_RNA       109,690,716       46.39       53.61 <t< td=""><td>3 MO RNA</td><td>50,716.752</td><td>46.26</td><td>53.74</td><td>98.66</td><td>95.85</td></t<>	3 MO RNA	50,716.752	46.26	53.74	98.66	95.85
3-       1_MO_RNA       49,042,662       46.46       53.54       98.67       95.82         3-1_NE_RNA       82,768,782       45.87       54.13       98.59       95.61         3-       2_MO_RNA       51,651,078       46.35       53.65       98.85       96.31         3-2_NE_RNA       87,883,904       46.60       53.40       98.59       95.59         3-       3_MO_RNA       54,955,000       46.32       53.68       98.84       96.34         3-NE_RNA       52,636,722       47.58       52.42       98.66       95.84         4-       1_MO_RNA       46,153,778       46.05       53.95       98.86       96.30         4-1_NE_RNA       53,096,182       46.73       53.27       98.67       95.93         4-       2_MO_RNA       42,770,800       46.44       53.57       98.83       96.33         4-2_NE_RNA       53,019,634       46.91       53.09       98.69       95.81         4-       3_MO_RNA       109,690,716       46.39       53.61       98.85       96.19         4-3_NE_RNA       47,729,898       46.34       53.66       98.71       95.92       5-         5-       1_MO_RNA	2-3 NE RNA	98.607.242	45.75	54.26	98.69	95.75
1_MO_RNA       49,042,662       46.46       53.54       98.67       95.82         3-1_NE_RNA       82,768,782       45.87       54.13       98.59       95.61         3-       -       -       -       -       -       -       -       -       -       -       -       95.82       95.61       -       -       -       -       -       -       95.61       -       -       -       -       -       -       -       -       95.61       -	3-					
3-1_NE_RNA       82,768,782       45.87       54.13       98.59       95.61         3-       3-       3-       3-       3-       3-       3-         2_MO_RNA       51,651,078       46.35       53.65       98.85       96.31         3-2_NE_RNA       87,883,904       46.60       53.40       98.59       95.59         3-       3_MO_RNA       54,955,000       46.32       53.68       98.84       96.34         3-3_NE_RNA       52,636,722       47.58       52.42       98.66       95.84         4-       -       -       -       -       -       -         1_MO_RNA       46,153,778       46.05       53.95       98.86       96.30         4-1_NE_RNA       53,996,182       46.73       53.27       98.86       96.33         4-2_MO_RNA       42,770,800       46.44       53.57       98.83       96.33         4-2_NE_RNA       53,019,634       46.91       53.09       98.69       95.81         4-       -       -       -       -       -       -         3_MO_RNA       109,690,716       46.39       53.61       98.85       96.19         4-3_NE_RNA       47,72	1 MO RNA	49.042.662	46.46	53.54	98.67	95.82
3-       3-       3-         2_MO_RNA       51,651,078       46.35       53.65       98.85       96.31         3-2_NE_RNA       87,883,904       46.60       53.40       98.59       95.59         3-       3_MO_RNA       54,955,000       46.32       53.68       98.84       96.34         3-3_NE_RNA       52,636,722       47.58       52.42       98.66       95.84         4-       -       -       -       -       -         1_MO_RNA       46,153,778       46.05       53.95       98.86       96.30         4-1_NE_RNA       53,996,182       46.73       53.27       98.67       95.93         4-       -       -       -       -       -       -         2_MO_RNA       42,770,800       46.44       53.57       98.83       96.33         4-2       -       -       -       -       -       -         3_MO_RNA       42,770,800       46.44       53.57       98.83       96.33         4-3       -       -       -       -       -       -       -         3_MO_RNA       109,690,716       46.39       53.61       98.85       96.19	3-1 NE RNA	82,768,782	45.87	54 13	98.59	95.61
2_MO_RNA       51,651,078       46.35       53.65       98.85       96.31         3-2_NE_RNA       87,883,904       46.60       53.40       98.59       95.59         3-       -       -       -       -       -         3_MO_RNA       54,955,000       46.32       53.68       98.84       96.34         3-3_NE_RNA       52,636,722       47.58       52.42       98.66       95.84         4-       -       -       -       -       -         1_MO_RNA       46,153,778       46.05       53.95       98.86       96.30         4-1_NE_RNA       53,996,182       46.73       53.27       98.67       95.93         4-       -	3-	02)/ 00)/ 02	10.07	01.10	20.02	20.01
3-2_NE_RNA 87,883,904 46.60 53.40 98.59 95.59 3- 3_MO_RNA 54,955,000 46.32 53.68 98.84 96.34 3-3_NE_RNA 52,636,722 47.58 52.42 98.66 95.84 4- 1_MO_RNA 46,153,778 46.05 53.95 98.86 96.30 4-1_NE_RNA 53,996,182 46.73 53.27 98.67 95.93 4- 2_MO_RNA 42,770,800 46.44 53.57 98.83 96.33 4-2_NE_RNA 53,019,634 46.91 53.09 98.69 95.81 4- 3_MO_RNA 109,690,716 46.39 53.61 98.85 96.19 4-3_NE_RNA 47,729,898 46.34 53.66 98.71 95.92 5- 1_MO_RNA 94,239,798 46.40 53.60 98.68 95.97 5-1_NE_RNA 48,834,504 46.25 53.75 98.78 96.07 5- 2_MO_RNA 99,231,268 46.44 53.56 98.74 96.03 5-2 NE_RNA 41 094 770 45.74 54.26 98.75 95.92	2 MO RNA	51,651,078	46.35	53 65	98 85	96.31
3-       3-       3-         3_MO_RNA       54,955,000       46.32       53.68       98.84       96.34         3-3_NE_RNA       52,636,722       47.58       52.42       98.66       95.84         4-       1_MO_RNA       46,153,778       46.05       53.95       98.86       96.30         4-1_NE_RNA       53,996,182       46.73       53.27       98.67       95.93         4-       2_MO_RNA       42,770,800       46.44       53.57       98.83       96.33         4-2.NE_RNA       53,019,634       46.91       53.09       98.69       95.81         4-       3_MO_RNA       109,690,716       46.39       53.61       98.85       96.19         4-3_NE_RNA       47,729,898       46.34       53.66       98.71       95.92       5-         1_MO_RNA       94,239,798       46.40       53.60       98.68       95.97         5-1_NE_RNA       48,834,504       46.25       53.75       98.78       96.07         5-       2_MO_RNA       99,231,268       46.44       53.56       98.74       96.03         5-2       NE_RNA       41.094,770       45.74       54.26       98.75       95.92 <td>3-2 NE RNA</td> <td>87,883,904</td> <td>46.60</td> <td>53 40</td> <td>98.59</td> <td>95.59</td>	3-2 NE RNA	87,883,904	46.60	53 40	98.59	95.59
3_MO_RNA       54,955,000       46.32       53.68       98.84       96.34         3-3_NE_RNA       52,636,722       47.58       52.42       98.66       95.84         4-       1_MO_RNA       46,153,778       46.05       53.95       98.86       96.30         4-1_NE_RNA       53,996,182       46.73       53.27       98.67       95.93         4-       -       -       -       -       -         2_MO_RNA       42,770,800       46.44       53.57       98.83       96.33         4-2_NE_RNA       53,019,634       46.91       53.09       98.69       95.81         4-       -       -       -       -       -       -         3_MO_RNA       109,690,716       46.39       53.61       98.85       96.19         4-3_NE_RNA       47,729,898       46.34       53.66       98.71       95.92         5-       -       -       -       -       -       -         1_MO_RNA       94,239,798       46.40       53.60       98.71       95.92         5-       -       -       -       -       -       -         1_MO_RNA       94,239,798       46.40       5	3-	07,000,001	10.00	00.10	20.02	20.02
3-3_NE_RNA       52,636,722       47.58       52.42       98.66       95.84         4-       1_MO_RNA       46,153,778       46.05       53.95       98.86       96.30         4-1_NE_RNA       53,996,182       46.73       53.27       98.67       95.93         4-       2_MO_RNA       42,770,800       46.44       53.57       98.83       96.33         4-2_NE_RNA       53,019,634       46.91       53.09       98.69       95.81         4-       -       -       -       -       -         3_MO_RNA       109,690,716       46.39       53.61       98.85       96.19         4-3_NE_RNA       47,729,898       46.34       53.66       98.71       95.92         5-       -       -       -       -       -         1_MO_RNA       94,239,798       46.40       53.60       98.68       95.97         5-1_NE_RNA       48,834,504       46.25       53.75       98.78       96.07         5-       -       -       -       -       -       -       -         46.40       53.56       98.74       96.03       -       -       -       -         5-2_NE_RNA	3 MO RNA	54,955,000	46.32	53 68	98 84	96.34
4-       1.00       52.12       90.00       90.01         4-       1_MO_RNA       46,153,778       46.05       53.95       98.86       96.30         4-1_NE_RNA       53,996,182       46.73       53.27       98.67       95.93         4-       2_MO_RNA       42,770,800       46.44       53.57       98.83       96.33         4-2_NE_RNA       53,019,634       46.91       53.09       98.69       95.81         4-       -       -       -       -       -         3_MO_RNA       109,690,716       46.39       53.61       98.85       96.19         4-3_NE_RNA       47,729,898       46.34       53.66       98.71       95.92         5-       -       -       -       -       -       -         1_MO_RNA       94,239,798       46.40       53.60       98.68       95.97       -         5-1_NE_RNA       48,834,504       46.25       53.75       98.78       96.07       -         5-       -       -       -       -       -       -       -         5-       -       -       -       -       -       -       -       -         5-	3-3 NE RNA	52 636 722	47.58	52 42	98.66	95.84
1_MO_RNA       46,153,778       46.05       53.95       98.86       96.30         4-1_NE_RNA       53,996,182       46.73       53.27       98.67       95.93         4-       -	4-	02,000,722	17.00	02.12	20.00	20.01
4-1_NE_RNA       53,996,182       46.73       53.27       98.67       95.93         4-       2_MO_RNA       42,770,800       46.44       53.57       98.83       96.33         4-2_NE_RNA       53,019,634       46.91       53.09       98.69       95.81         4-       3_MO_RNA       109,690,716       46.39       53.61       98.85       96.19         4-3_NE_RNA       47,729,898       46.34       53.66       98.71       95.92         5-       5-       5-       53.60       98.68       95.97         5-1_NE_RNA       48,834,504       46.25       53.75       98.78       96.07         5-       2_MO_RNA       99,231,268       46.44       53.56       98.74       96.03         5-2       NE_RNA       41.094.770       45.74       54.26       98.75       95.92	1 MO RNA	46,153,778	46.05	53 95	98 86	96.30
4-       10.70       50.27       90.87       90.87         2_MO_RNA       42,770,800       46.44       53.57       98.83       96.33         4-2_NE_RNA       53,019,634       46.91       53.09       98.69       95.81         4-       -       -       -       -       -         3_MO_RNA       109,690,716       46.39       53.61       98.85       96.19         4-3_NE_RNA       47,729,898       46.34       53.66       98.71       95.92         5-       -       -       -       -       -         1_MO_RNA       94,239,798       46.40       53.60       98.68       95.97         5-1_NE_RNA       48,834,504       46.25       53.75       98.78       96.07         5-       -       -       -       -       -       -         2_MO_RNA       99,231,268       46.44       53.56       98.74       96.03         5-2       NE_RNA       41.094.770       45.74       54.26       98.75       95.92	4-1 NE RNA	53 996 182	46 73	53 27	98.67	95.93
2_MO_RNA       42,770,800       46.44       53.57       98.83       96.33         4-2_NE_RNA       53,019,634       46.91       53.09       98.69       95.81         4-       -       -       -       -       -       -         3_MO_RNA       109,690,716       46.39       53.61       98.85       96.19         4-3_NE_RNA       47,729,898       46.34       53.66       98.71       95.92         5-       -       -       -       -       -       -         1_MO_RNA       94,239,798       46.40       53.60       98.68       95.97         5-1_NE_RNA       48,834,504       46.25       53.75       98.78       96.07         5-       -       -       -       -       -       -         46.40       53.66       98.74       96.07       -       -       -         5-       -       -       -       -       -       -       -         1_MO_RNA       94,239,798       46.40       53.60       98.78       96.07       -         5-       -       -       -       -       -       -       -         2_MO_RNA       99,231,268 <td>4-</td> <td>00,770,102</td> <td>10.70</td> <td>00.27</td> <td>20.07</td> <td>20.20</td>	4-	00,770,102	10.70	00.27	20.07	20.20
4-2_NE_RNA 53,019,634 46.91 53.09 98.69 95.81 4- 3_MO_RNA 109,690,716 46.39 53.61 98.85 96.19 4-3_NE_RNA 47,729,898 46.34 53.66 98.71 95.92 5- 1_MO_RNA 94,239,798 46.40 53.60 98.68 95.97 5-1_NE_RNA 48,834,504 46.25 53.75 98.78 96.07 5- 2_MO_RNA 99,231,268 46.44 53.56 98.74 96.03 5-2 NE_RNA 41.094.770 45.74 54.26 98.75 95.92	2 MO RNA	42 770 800	46 44	53 57	98 83	96.33
4-       3_MO_RNA       109,690,716       46.39       53.61       98.85       96.19         4-3_NE_RNA       47,729,898       46.34       53.66       98.71       95.92         5-       5-       5-       5-       5-       5-         1_MO_RNA       94,239,798       46.40       53.60       98.68       95.97         5-1_NE_RNA       48,834,504       46.25       53.75       98.78       96.07         5-       5-       5-       5-       5-       5-       5-         2_MO_RNA       99,231,268       46.44       53.56       98.74       96.03         5-2       NE_RNA       41.094.770       45.74       54.26       98.75       95.92	4-2 NE RNA	53 019 634	46 91	53.09	98.69	95.81
3_MO_RNA 109,690,716 46.39 53.61 98.85 96.19 4-3_NE_RNA 47,729,898 46.34 53.66 98.71 95.92 5- 1_MO_RNA 94,239,798 46.40 53.60 98.68 95.97 5-1_NE_RNA 48,834,504 46.25 53.75 98.78 96.07 5- 2_MO_RNA 99,231,268 46.44 53.56 98.74 96.03 5-2 NE_RNA 41.094.770 45.74 54.26 98.75 95.92	4-	00,017,001	10.71	00.07	20.02	20.01
4-3_NE_RNA 47,729,898 46.34 53.66 98.71 95.92 5- 1_MO_RNA 94,239,798 46.40 53.60 98.68 95.97 5-1_NE_RNA 48,834,504 46.25 53.75 98.78 96.07 5- 2_MO_RNA 99,231,268 46.44 53.56 98.74 96.03 5-2 NE_RNA 41.094.770 45.74 54.26 98.75 95.92	3 MO RNA	109,690 716	46.39	53 61	98 85	96 19
5-       1_MO_RNA       94,239,798       46.40       53.60       98.68       95.97         5-1_NE_RNA       48,834,504       46.25       53.75       98.78       96.07         5-       2_MO_RNA       99,231,268       46.44       53.56       98.74       96.03         5-2       NE_RNA       41.094.770       45.74       54.26       98.75       95.92	4-3 NE RNA	47,729,898	46.34	53.66	98 71	95 92
1_MO_RNA       94,239,798       46.40       53.60       98.68       95.97         5-1_NE_RNA       48,834,504       46.25       53.75       98.78       96.07         5-       -       -       -       -       -       -         2_MO_RNA       99,231,268       46.44       53.56       98.74       96.03         5-2       NE       RNA       41.094.770       45.74       54.26       98.75       95.92	5-	<u>.,,,,,,,,,,</u> ,,,,,,,,,,,,,,,,,,,,,,,,,,	10.01	00.00	20.7 I	, U, JZ
5-1_NE_RNA 48,834,504 46.25 53.75 98.78 96.07 5- 2_MO_RNA 99,231,268 46.44 53.56 98.74 96.03 5-2 NE_RNA 41,094,770 45.74 54.26 98.75 95.92	1 MO RNA	94,239,798	46 40	53 60	98 68	95 97
5- 2_MO_RNA 99,231,268 46.44 53.56 98.74 96.03 5-2 NE RNA 41.094.770 45.74 54.26 98.75 95.92	5-1 NE RNA	48,834 504	46 25	53 75	98 78	96.07
2_MO_RNA 99,231,268 46.44 53.56 98.74 96.03 5-2 NE RNA 41 094 770 45 74 54 26 98 75 95 92	5-	10,001,001	10.20	00.70	20.70	20.07
5-2 NE RNA 41 094 770 45 74 54 26 98 75 95 92	2 MO RNA	99 231 268	46 44	53 56	98 74	96.03
-1/2 $-1/2$ $-1/2$ $-1/2$ $-1/2$ $-1/2$ $-1/2$ $-1/2$ $-1/2$	5-2 NF RNA	41 094 770	45.74	54 26	98 75	95 92
5-	5-	±1,07±,770	-10.7 I	07.20	20.70	JU.JL
3 MO RNA 88 308 474 46 46 53 55 98 68 95 94	3 MO RNA	88.308 474	46 46	53 55	98 68	95 94

**Supplementary Table S10.** Resume of read counts and statistics for each sequenced sample in the NovaSeq run.

**Supplementary Table S11.** List of accession numbers retrieved from NCBI following integration with the custom database (see Methods section) for the detection of viruses, viroids and phytoplasmas.

Accession	Virus / Viroid / Phitoplasma name				
NC_022002.1	Grapevine red blotch-associated virus	NC_002692.1	Tomato mosaic virus	NC_011535.1	Grapevine Algerian latent virus
NC_015784.2	Grapevine vein-clearing virus	LT608395.1	Artichoke italian latent virus RNA 1	AY500881.1	Petunia asteroid mosaic virus
	Grapevine roditis leaf discoloration-associated				
NC_027131.1	virus	LT608396.1	Artichoke italian latent virus RNA 2	NC_031692.1	Grapevine asteroid mosaic-associated virus
NC_035939.1	Grapevine Cabernet Sauvignon reovirus NSS	NC_005289.1	Broad bean wilt virus 1 RNA 1	NC_012484.1	Grapevine Syrah Virus-1
NC_035938.1	Grapevine Cabernet Sauvignon reovirus P7	NC_005290.1	Broad bean wilt virus 1 RNA 2	KM491303.1	Grapevine Red Globe virus
NC_035937.1	Grapevine Cabernet Sauvignon reovirus P6	NC_006057.1	Arabis mosaic virus RNA 1	NC_034205.1	Grapevine rupestris vein feathering virus
NC_035936.1	Grapevine Cabernet Sauvignon reovirus P4	NC_006056.1	Arabis mosaic virus RNA 2	NC_003347.1	Grapevine fleck virus
NC_035935.1	Grapevine Cabernet Sauvignon reovirus P1	NC_015414.1	Cherry leaf roll virus RNA 1	FJ915122.1	Blackberry virus S
	Grapevine Cabernet Sauvignon reovirus segment				
NC_027802.1	9	NC_015415.1	Cherry leaf roll virus RNA 2	NC_003739.1	Raspberry bushy dwarf virus RNA 1
NC_027816.1	Grapevine Cabernet Sauvignon reovirus P2	U20622.1	Blueberry leaf mottle virus RNA1	NC_003740.1	Raspberry bushy dwarf virus RNA 2
NC_027810.1	Grapevine Cabernet Sauvignon reovirus P10a	U20621.1	Blueberry leaf mottle virus RNA2	NC_011187.1	Rubus chlorotic mottle virus
NC_027809.1	Grapevine Cabernet Sauvignon reovirus P5	NC_015492.1	Grapevine Bulgarian latent virus RNA 1	KX949574.1	Southern tomato virus
NC_027808.1	Grapevine Cabernet Sauvignon reovirus P3a	NC_015493.1	Grapevine Bulgarian latent virus RNA 2	KX962563.1	Grapevine fabavirus RNA 1
NC_019493.1	Grapevine endophyte endornavirus	NC_018383.1	Grapevine Anatolian ringspot virus RNA 1	KX962564.1	Grapevine fabavirus RNA 2
NC_011705.1	Raphanus sativus cryptic virus 3 segment 1	NC_018384.1	Grapevine Anatolian ringspot virus RNA 2	HG939487.1	Tomato black ring virus RNA 1
NC_011706.1	Raphanus sativus cryptic virus 3 segment 2	NC_017939.1	Grapevine deformation virus RNA1	KX977561.1	Tomato black ring virus RNA 2
S63913.1	Beet cryptic virus 3	NC_017938.1	Grapevine deformation virus RNA2	KX645875.2	Grapevine enamovirus-1
NC_002050.1	Tomato spotted wilt virus RNA M	NC_003622.1	Grapevine chrome mosaic virus RNA 1	GQ845002.2	Sowbane mosaic virus
NC_002051.1	Tomato spotted wilt virus RNA S	NC_003621.1	Grapevine chrome mosaic virus RNA 2	JX658571.1	Grapevine partitivirus
NC_002052.1	Tomato spotted wilt virus RNA L	NC_003615.1	Grapevine fanleaf virus RNA 1	NC_035203.1	Grapevine virus T
NC_029783.1	Grapevine leafroll-associated virus 13	NC_003623.1	Grapevine fanleaf virus RNA 2	MF781081.1	Grapevine virus G
NC_007448.1	Grapevine leafroll-associated virus 2	NC_034214.1	Peach rosette mosaic virus RNA 1	MF521889.1	Grapevine virus H
NC_004667.1	Grapevine leafroll-associated virus 3	NC_034215.1	Peach rosette mosaic virus RNA 2	NC_037058.1	Grapevine virus I
NC_016509.1	Grapevine leafroll-associated virus 1	NC_005266.1	Raspberry ringspot virus RNA 1	MG637048.1	Grapevine virus J
NC_016436.1	Grapevine leafroll-associated virus 7	NC_005267.1	Raspberry ringspot virus RNA 2	NC_035480.1	Wild vitis virus 1
NC_016081.1	Grapevine leafroll-associated virus 5	NC_005097.1	Tobacco ringspot virus RNA 1	KX950822.1	Grapevine geminivirus A
NC_011702.1	Grapevine leafroll-associated virus 10	NC_005096.1	Tobacco ringspot virus RNA 2	KJ955447.1	Temperate fruit decay-associated virus
NC_016417.1	Grapevine leafroll-associated virus 6	NC_003840.1	Tomato ringspot virus RNA 1	NC_015784.2	Grapevine vein-clearing virus
NC_016416.1	Grapevine leafroll-associated virus 4	NC_003839.2	Tomato ringspot virus RNA 2	MF781082.1	Grapevine badnavirus 1
NC_011620.1	Potato virus X	NC_006964.1	Strawberry latent ringspot virus RNA 1	NC_001920.1	Grapevine yellow speckle viroid 1
NC_001948.1	Rupestris stem pitting associated virus-1	NC_006965.1	Strawberry latent ringspot virus RNA 2	NC_003612.1	Grapevine yellow speckle viroid 2
NC_015220.1	Grapevine berry inner necrosis virus	NC_001495.1	Alfalfa mosaic virus RNA 1	NC_003553.1	Australian grapevine viroid
NC_015782.1	Grapevine Pinot gris virus	NC_002024.2	Alfalfa mosaic virus RNA 2	NC_001351.1	Hop stunt viroid

NC_003604.2	Grapevine virus A	NC_002025.1	Alfalfa mosaic virus RNA 3	NC_001464.1	Citrus exocortis viroid
NC_003602.1	Grapevine virus B	NC_002034.1	Cucumber mosaic virus RNA 1	JQ046414.1	Grapevine yellows phytoplasma 16S
MF774336.1	Grapevine virus D	NC_002035.1	Cucumber mosaic virus RNA 2	JQ900580.1	Grapevine yellows phytoplasma rpl22
NC_011106.1	Grapevine virus E	NC_001440.1	Cucumber mosaic virus RNA 3	AF385627.1	Flavescence doree phytoplasma rpl22
NC_018458.1	Grapevine virus F	AY590305.1	Grapevine angular mosaic virus	AF396951.1	Flavescence doree phytoplasma rps3
NC_003397.1	Bean common mosaic virus	NC_001265.2	Carnation mottle virus	JQ181540.1	Bois noir' phytoplasma 16S
NC_001367.1	Tobacco mosaic virus	NC_003487.1	Tobacco necrosis virus D		

**Figure S1.** PCA analysis of EcoPlate<sup>TM</sup> data. Distinctions between cultivar 'Moscato' (MO) and 'Nebbiolo' (NE) as revealed by PCA over the time of plate incubation from day 1 (h24) to 5 (h120). Numbers reported after the name of the cultivar (NE: 'Nebbiolo', MO: 'Moscato') refer to the treatment-type as following stated: 1 = Inoculated untreated control (CTRL), 2 = Acibenzolar-S-methyl (AcS-Mt), 3 = Potassium Phosphonate (K-Pho), and 4 = Laminarin (Lam).



**Figure S2.** PCA analysis of EcoPlate<sup>TM</sup> data. Distinctions between cultivar 'Moscato' (MO) and 'Nebbiolo' (NE) compared within each treatment as revealed by PCA over the time of plate incubation from day 1 (h24) to 5 (h120). (a) Inoculated untreated control (CTRL), (b) Acibenzolar-S-methyl (AcS-Mt), (c) Potassium Phosphonate (K-Pho), and (d) Laminarin (Lam).



**Figure S3.** Average Well Colour Development (AWCD) in Biolog EcoPlates<sup>TM</sup> inoculated with 'Moscato' (a) and 'Nebbiolo' (b) leaf sample extracts. Values are calculated on the base of absorbance readings taken at 590 nm for each plate over a time course of 5 days (h24 to h120). Data represent mean ± standard deviation of three biological replicates. Inoculated untreated control (CTRL), Acibenzolar-S-methyl (AcS-Mt), Potassium Phosphonate (K-Pho), and Laminarin (Lam).



**Figure S4.** Rarefaction OTU curves obtained for each biological replicate of 'Moscato' (MO, panels from a to d) and 'Nebbiolo' (NE, panels from e to h) leaf samples collected from untreated (CTRL) and treated (AcS-Mt = acibenzolar-S-methyl, K-Pho = Potassium Phosphonate, Lam = Laminarin) plants infected by powdery mildew.



**Figure S5.** Agarose gel electrophoretic analysis of DNA fragments amplified by multiplex RT-PCR for detection of virus (**a** and **b**, Gambino 2015), viroids (**b**, Hajizadeh et al., 2012) and Grapevine pinot gris virus, GPGV (**c**, Glasa et al., 2014) in 'Moscato' and 'Nebbiolo'. CTRL - Inoculated untreated control, AcS-Mt - Acibenzolar-S-methyl, K-Pho - Potassium Phosphonate, LAM – Laminarin. NE: 'Nebbiolo', MO: 'Moscato'.

