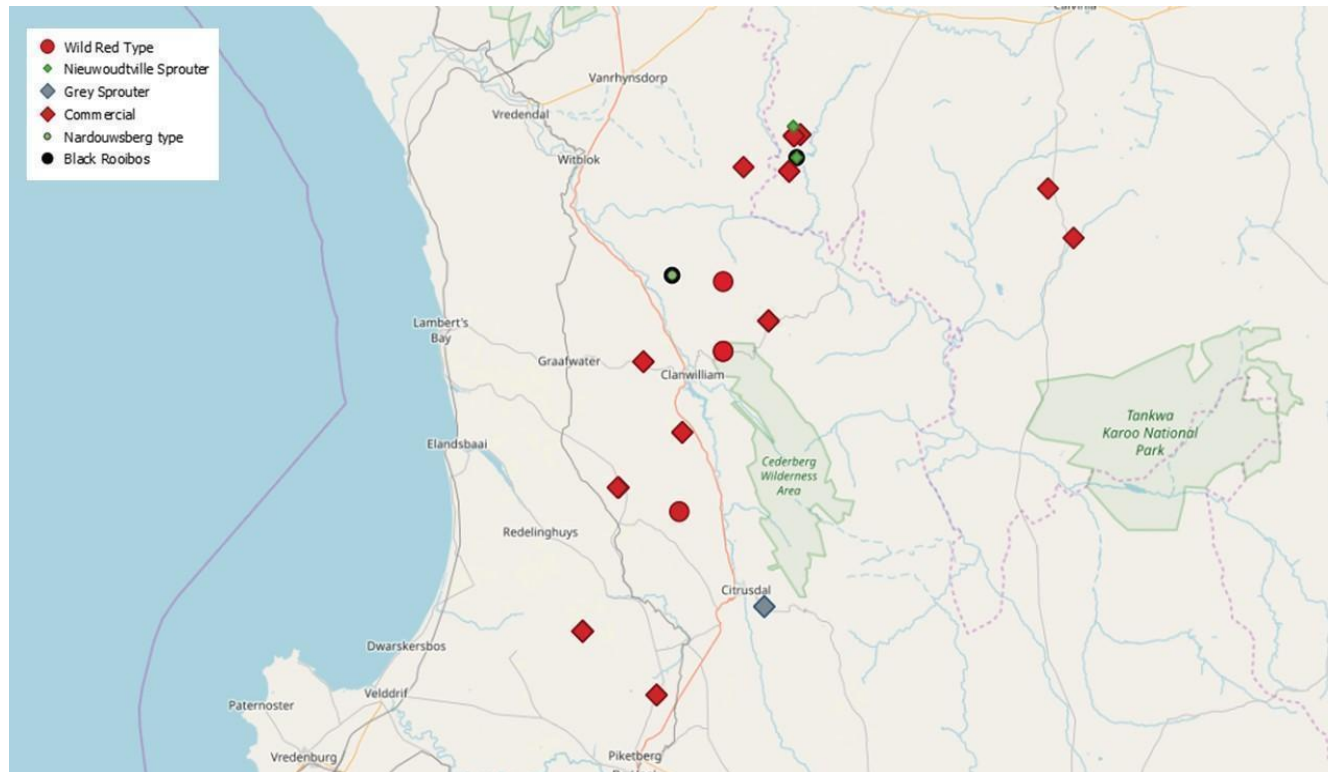


Supplementary Materials: Transcriptomics of the Rooibos (*Aspalathus linearis*) Species Complex

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Supplementary Figure S1: Distribution of the collected rooibos ecotypes sampled for this study.

Supplementary Table S1: Number of overlapping orthogroups predicted by OrthoFinder.

	number of sequences	A	B	C	D	<i>L. angustifolius</i>	<i>L. japonicus</i>	<i>M. truncatula</i>	<i>A. thaliana</i>	<i>O. sativa</i>
A	85234	29535	20486	19144	19629	12744	13471	13333	11429	10943
B	91301	20486	28650	19545	19755	12819	13546	13439	11485	10992
C	74767	19144	19545	26334	18968	12823	13434	13360	11489	10937
D	79234	19629	19755	18968	26912	12604	13296	13187	11357	10873
<i>L. angustifolius</i>	33076	12744	12819	12823	12604	14308	12712	13185	11646	10777
<i>L. japonicus</i>	48105	13471	13546	13434	13296	12712	16268	13719	11564	10932
<i>M. truncatula</i>	57585	13333	13439	13360	13187	13185	13719	15828	11987	11229
<i>A. thaliana</i>	35386	11429	11485	11489	11357	11646	11564	11987	12665	10685
<i>O. sativa</i>	66338	10943	10992	10937	10873	10777	10932	11229	10685	12526