

Development of SSR markers for five mangrove species using next-eneration sequencing with preliminary results of analyses of genetic diversity

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Abstract	
Authors	Y Shinmura, K Takayama, WK Shan, EL Webb, T Asakawa, B Adjie, ...
Journal Name	VLIZ Special Publication, 2012
Publish Year	2012
Citation	(not set)
Url	https://scholar.google.com/scholar?q=+intitle:"Development of SSR markers for five mangrove species using next-eneration sequencing with preliminary results of analyses of genetic diversity"
Author	Dr.rer.nat. ERWIN RIYANTO ARDLI, M.Sc.