Supporting Information

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Isolation and Characterization of Glycosidic Tyrosinase

Inhibitors from Typhonium giganteum Rhizomes

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Single Mass Analysis Tolerance = 5.0 PPM / DBE: min = -1.5, max = 50.0 Element prediction: Off Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions 236 formula(e) evaluated with 1 results within limits (up to 50 closest results for each mass) Elements Used: C: 15-15 H: 0-100 N: 0-10 O: 0-10 Na: 0-1

1:	TOF	MS	ES+	

100	301.1 300	404 31 305	2.1919 	320.8 315	695 32 320	25.221 777 325	3 330	338. 335	3471 	345	351.10 	064 354 355	.2838 360	367.0 365	0668 / 370	71.2921	38 380	4.3495 	393 	2987 395	400	1.66e+00 406.3296 +++++ m/z 405	4
Minim Maxim	um: um:			20.0	5.0	-1 50	. 5																
Mass		Calc.	Mass	mDa	PPM	DE	BE	i-FIT	Nor	m C	Conf(%)	Formu	la										
351.1	064	351.1	056	0.8	2.3	5.	5	403.9	n/a	n	ı/a	C15 H	120 08	Na									

Figure S1: HR-ESI-MS spectrum of 1



Figure S2: IR spectrum of 1



Figure S3: UV spectrum of 1 in MeOH



Figure S4: ¹H NMR spectrum (400 MHz) of 1 in CD₃OD



Figure S5: ¹³C NMR spectrum (100 MHz) of 1 in CD₃OD



Figure S6: DEPT 135 spectrum of 1 in CD₃OD



Figure S7: HSQC spectrum of 1 in CD₃OD



Figure S8: ¹H-¹H COSY spectrum of 1 in CD₃OD



Figure S9: HMBC spectrum of 1 in CD₃OD



Figure S10: NOESY spectrum of 1 in CD₃OD



Figure S11: ¹H NMR spectrum (400 MHz) of 1a in CD₃OD



Figure S12: ¹³C NMR spectrum (100 MHz) of 1a in CD₃OD