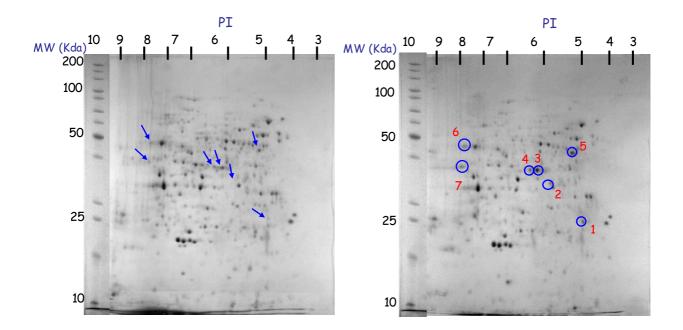
## Metabolic networking in *Brunfelsia calycina* petals during flower opening

Ayelet Bar-Akiva<sup>\*1</sup>, Rinat Ovadia<sup>\*1</sup>, Ilana Rogachev<sup>\*2</sup>, Carmiya Bar-Or<sup>1</sup>, Einat Bar<sup>3</sup>, Zohar Freiman<sup>1</sup>, Ada Nissim-Levi<sup>1</sup>, Natan Gollop<sup>4</sup>, Efraim Lewinsohn<sup>3</sup>, Asaph Aharoni<sup>2</sup>, David Weiss<sup>5</sup>, Hinanit Koltai<sup>1</sup> and Michal Oren-Shamir<sup>\*\*1</sup>



**Figure S1**. 2-D gels of total protein extracts from Brunfelsia flowers at D0 (left) and D2 (right) after opening. The first IEF dimension was on 13 cm dry strips (pH3-10), and the second dimension was on 12% acrylamide SDS-PAGE. Each gel is a computed average of five 2-D gels. The seven marked protein spots showed at least 3- fold increased intensity after Coomassie staining.