

The results shown are the average \pm standard deviation of the results of three experiments where each timepoint was analyzed in triplicate. (ND = sample not analyzed.)

To compare the results obtained using full scan versus selected ion monitoring (SIM), the reaction of aniline and benzylbromide was repeated. Data were collected using full scan mode and the selected ion monitoring of the key ions of aniline and dibenzylphenylamine. The results of the comparison are shown in the Figure 3.

SUMMARY

- Direct analysis of a reaction mixture by a modified APCI source with the ASAP probe provided unequivocal reaction identification.
- Easy-to-use and reliable mass analysis system
- Results are generated in less than 30 seconds

TIME	CYCLOHEXYLAMINE REACTION		ANILINE REACTION
	ROOM TEMPERATURE	ICE BATH	ROOM TEMPERATURE
1	95.91 \pm 2.20	0.00 \pm 0.00	1.66 \pm 1.46
5	99.44 \pm 0.13	74.53 \pm 10.39	11.11 \pm 2.98
10	99.60 \pm 0.09	95.38 \pm 1.79	18.55 \pm 2.52
30	98.91 \pm 0.59	97.82 \pm 1.03	60.66 \pm 8.13
60	99.47 \pm 0.23	98.85 \pm 1.22	73.24 \pm 5.29
120	99.68 \pm 0.21	99.70 \pm 0.03	82.15 \pm 14.87
180	99.76 \pm 0.04	ND	89.81 \pm 5.95

Table 1: Aniline and cyclohexylamine with benzylbromide analyzed at time points by ASAP enabled APCI source on CMS.

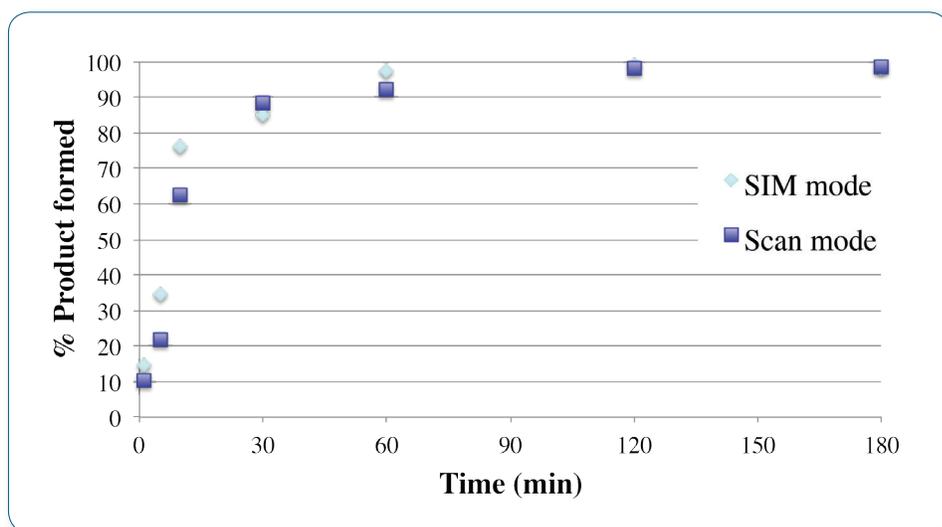


Figure 3: The analysis indicates that there is close correspondence between the results obtained using either full scan or SIM monitoring.

REFERENCES

¹ Jason L. Moore, Stephen M. Taylor and Vadim A. Soloshonok, *Archive for Organic Chemistry*, 2005 Vol 6, p 287-292

Charles N. McEwen,* Richard G. McKay, and Barbara S. Larsen, *Anal. Chem.* 2005, 77, 7826-7831

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