

SUPPLEMENTAL***Hydrophobic Treatment Enabling Analysis of Wettable Surfaces using a Liquid Microjunction Surface Sampling Probe/Electrospray Ionization-Mass Spectrometry System*****Matthew J. Walworth,^{1,2} Joseph J. Stankovich,¹ and Gary J. Van Berkel,^{1*}****Michael Schulz³ and Susanne Minarik³****Judy Nichols⁴ and Eike Reich⁵**

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Submitted for consideration for publication in
Analytical Chemistry

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Figure S1. Photographs showing the LMJ-SSP setup on LTQ mass spectrometer. (a) Closeup view of LMJ-SSP and atmospheric pressure interface region of mass spectrometer. (b) Wide angle view of complete instrumental setup.

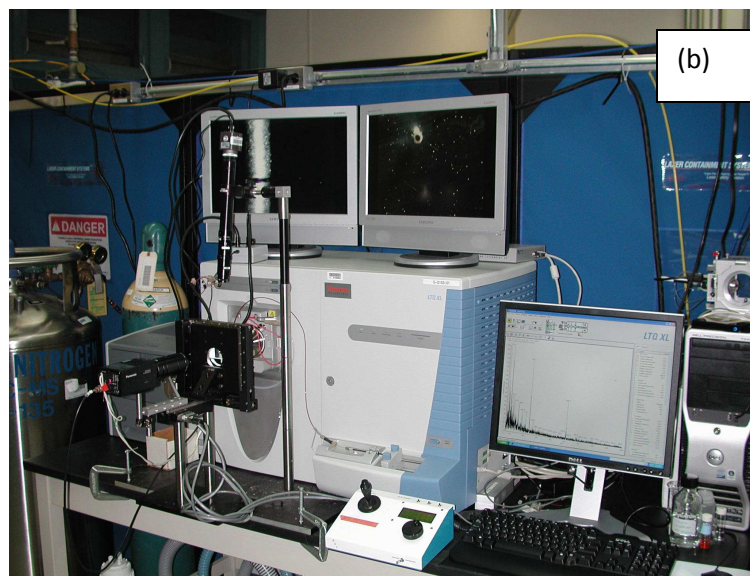
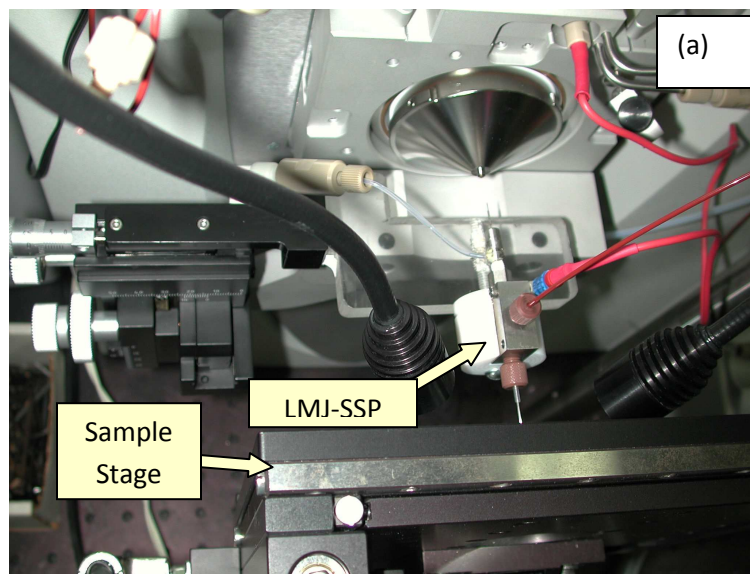


Figure S2. Multiple absorbent surface types were successfully treated to created hydrophobic character needed for successful LMJ-SSP analysis. Beading of water on surface is one test used to gauge the quality of surface treatment.

Hydrophobic Surface Treatment

- Water beads on surface
 - test for successful surface treatment



Off the shelf
hydrophobic Reverse-
Phase HPTLC plate
No Treatment

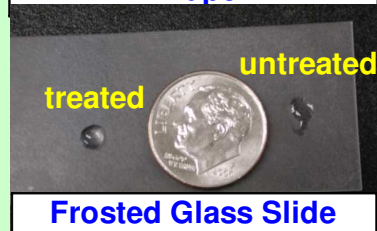
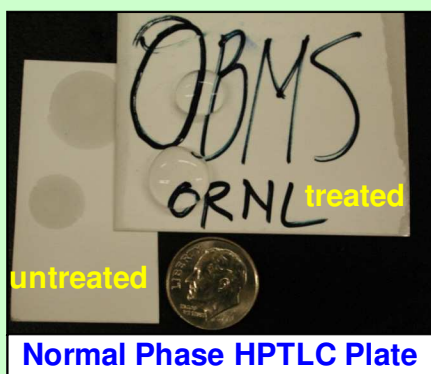


Figure S3. (a) Silicone treated kimwipe marked prior to treatment with black sharpie marker lines. (b) Extracted ion current profile for the black dye component (m/z 372) obtained during LMJ-SSP scan at $100 \mu\text{m s}^{-1}$ along treated kimwipe. (c) Averaged mass spectrum acquired over the line labeled (c) on the kimwipe. The extraction solvent was composed of 60/40 methanol/water (v/v). Data was acquired in full scan mode (m/z 150 – 2000).

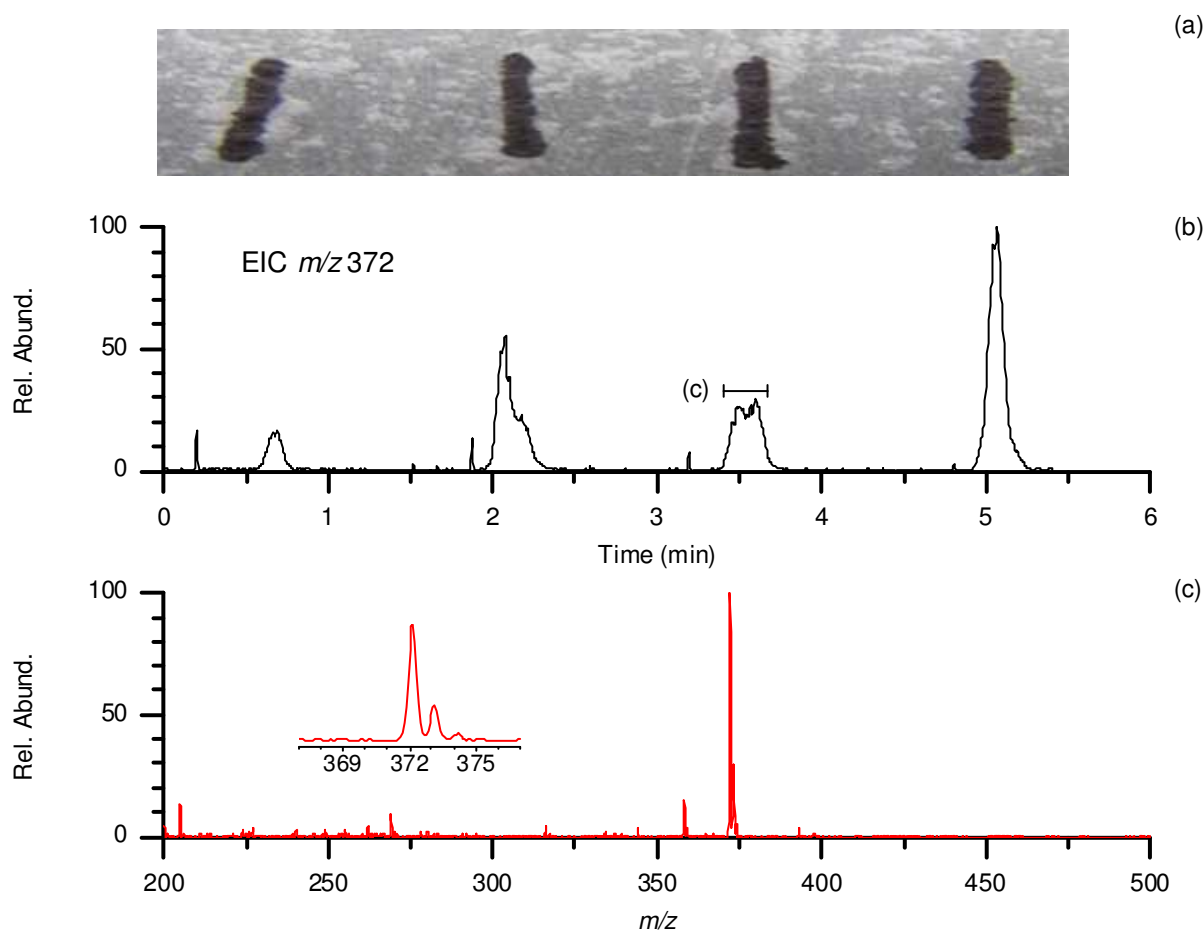


Figure S4. Representative mass spectrum of silicone polymer signal that is observed when treated surfaces are not prepared properly.

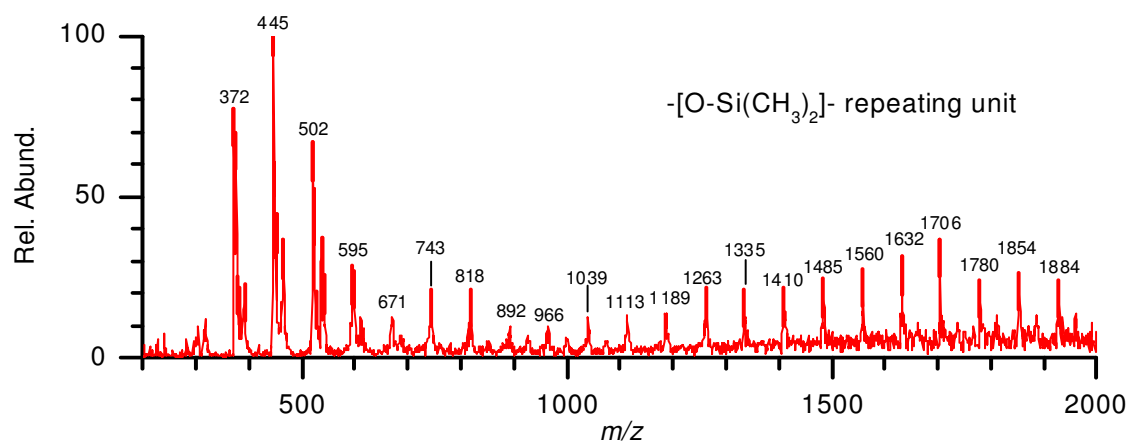


Figure S5. Peptide ID Information

Amino acids in black font and underlined was identified by MS/MS database searching. Amino acids in red font were not identified.

BSA Digest

Proteochrome HPTLC cellulose plate

MKWVTFISLLLLFSSAYSSRGVFRRDTHKSEIAHRFKDLGEEHFKGLVLIAFSQYLQQCPF
DEHVKLNELTEFAKTCVADESHAGCEKSLHTLFGDELCKVASLRETYGDMADCCEKQE
PERNECFLSHKDDSPDLPKLKPDNTLCDEFKADEKKFWGKYLYEIARRHPYFYAPELLY
YANKYNGVFQECCQAEDKGACLLPKIETMREKVLTSSARQRLRCASIQKFGERALKAWS
VARLSQKFPKAEFVEVTKLVTDLTKVHKECCHGDLLECADDRADLAKYICDNQDTISSK
LKECCDKPLLEKSHCIAEVEKDAIPENLPPLTADFAEDKDVCKNYQEAKDAFLGSFLYEY
SRRHPEYAVSVLLRLAKEYEATLEECCAKDDPHACYSTVFDKLKHLVDEPQNLIKQNCD
QFEKLGEYGFQNALIVRYTRKVPQVSTPTLVEVSRSLGKVGTRCCTKPESERMPCTEDYL
SLILNRLCVLHEKTPVSEKVTKCTESLVNRRPCFSALTPDETYVPKAFDEKLLFTFHADIC
TLPDTEKQIKKQTALVELLKHKPKATEEQLKTVMENFVAFVDKCCAADDKEACFAVEGP
KLVVSTQTALA

Sequence Coverage 40.5%

Proteochrome HPTLC silica gel 60 plate

MKWVTFISLLLLFSSAYSSRGVFRRDTHKSEIAHRFKDLGEEHFKGLVLIAFSQYLQQCPF
DEHVKLNELTEFAKTCVADESHAGCEKSLHTLFGDELCKVASLRETYGDMADCCEKQE
PERNECFLSHKDDSPDLPKLKPDNTLCDEFKADEKKFWGKYLYEIARRHPYFYAPELLY
YANKYNGVFQECCQAEDKGACLLPKIETMREKVLTSSARQRLRCASIQKFGERALKAWS
VARLSQKFPKAEFVEVTKLVTDLTKVHKECCHGDLLECADDRADLAKYICDNQDTISSK
LKECCDKPLLEKSHCIAEVEKDAIPENLPPLTADFAEDKDVCKNYQEAKDAFLGSFLYEY
SRRHPEYAVSVLLRLAKEYEATLEECCAKDDPHACYSTVFDKLKHLVDEPQNLIKQNCD
QFEKLGEYGFQNALIVRYTRKVPQVSTPTLVEVSRSLGKVGTRCCTKPESERMPCTEDY
LSLILNRLCVLHEKTPVSEKVTKCTESLVNRRPCFSALTPDETYVPKAFDEKLLFTFHADI
CTLPDTEKQIKKQTALVELLKHKPKATEEQLKTVMENFVAFVDKCCAADDKEACFAVEG
PKLVVSTQTAL

Sequence Coverage 60.5%

Lysosyme Digest

Proteochrome HPTLC cellulose plate

KVFGRCELAAAMKRHGLDNYRGYSLGNWVCAAKFESNFNTQA
TNRNTDGSTDYGILQINSRWWCNDGRTPGSR**NLCNIPCSALLSS**
DITASVNCAKKIVSDGNGMNAWVAWR**NRCK**GTDVQAWIR**GCRL**

Sequence Coverage 75.2%

Proteochrome HPTLC silica gel 60 plate

KVFGRCELAAAMK**R**HGLDNYRGYSLGNWVCAAKFESNFNTQA
TNRNTDGSTDYGILQINSRWWCNDGR**TPGSR**NLCNIPCSALLSS
DITASVNCAK**K**IVSDGNGMNAWVAWR**NRCK**GTDVQAWIR**GCRL**

Sequence Coverage 88.4%

Myoglobin Digest

Proteochrome HPTLC cellulose plate

GLSDGEWQQV LNVWGKVEAD IAGHGQEVLI RLFTGHPETL
EKFDKFKHLKTEAEMKASEDLK**K**HGTVVLT ALGGILK**KKG HHEAELKPLA**
QSHATKHKIP IKYLEFISDA IIVVLHSHKHPGDFGADAQGA MTKALELFRN
DIAAKYKELG FQG

Sequence Coverage 98.0%

Proteochrome HPTLC silica gel 60 plate

GLSDGEWQQV LNVWGKVEAD IAGHGQEVLI RLFTGHPETL
EKFDKFKHLKTEAEMKASEDLK**K**HGTVVLT ALGGILK**KKG HHEAELKPLA**
QSHATKHKIP IKYLEFISDA IIVVLHSHKHPGDFGADAQGA MTKALELFRN
DIAAKYKELG FQG

Sequence Coverage 85.6%

Beta Casein Digest

Proteochrome HPTLC cellulose plate

MKVLILACLVALALARELEELNVPGEIVESLSSEESITRINKKIEKFQSEEQQQTEDELQD
KIHFAQTQSLVYPFPGPIPNLQNIPLTQTPVVVPPFLQPEVMGVSKVKEAMAPKHKE
MPFPKYPVEPFOTESQSLTLTDVENLHLPLLLQSWMHQPHQLPPTVMFPPQSVLSLSQS
KVLPVPQKAVPYPQORDMPIQAFLLYQEPVLGPVRGPFPIIV

Sequence Coverage 22.3%

Proteochrome HPTLC silica gel 60 plate

MKVLILACLVALALARELEELNVPGEIVESLSSEESITRINKKIEKFQSEEQQQTEDELQD
KIHFAQTQSLVYPFPGPIPNLQNIPLTQTPVVVPPFLQPEVMGVSKVKEAMAPKHKE
MPFPKYPVEPFOTESQSLTLTDVENLHLPLLLQSWMHQPHQLPPTVMFPPQSVLSLSQS
KVLPVPQKAVPYPQORDMPIQAFLLYQEPVLGPVRGPFPIIV

Sequence Coverage 29.5%

Bovine Cytochrome C Digest

Proteochrome HPTLC cellulose plate

GDVEKGKKIF VQKCAQCHTV EKGGKHKTGP NLHGLFGRKT GQAPGFSYTD
ANKNGKITWGEETLMEYLEN PKKYIPGTKM IFAGIKKKE REDLIAYLKK ATNE

Sequence Coverage 90.4%

Proteochrome HPTLC silica gel 60 plate

GDVEKGKKIF VQKCAQCHTV EKGGKHKTGP NLHGLFGRKT GQAPGFSYTD
ANKNGKITWGEETLMEYLEN PKKYIPGTKM IFAGIKKKE REDLIAYLKK ATNE

Sequence Coverage 89.4%