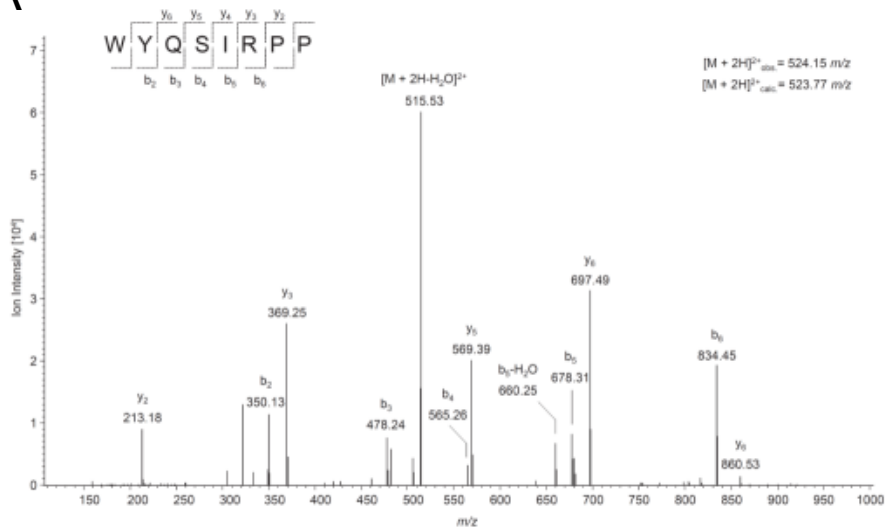
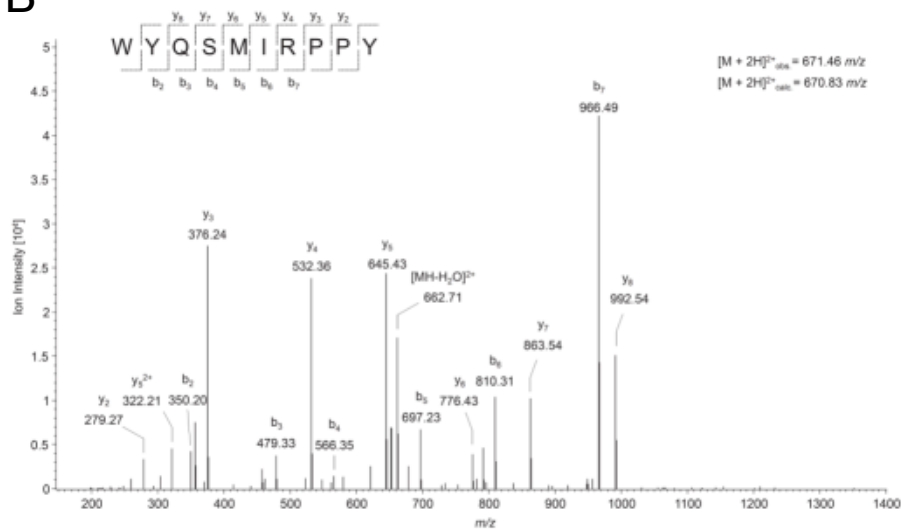


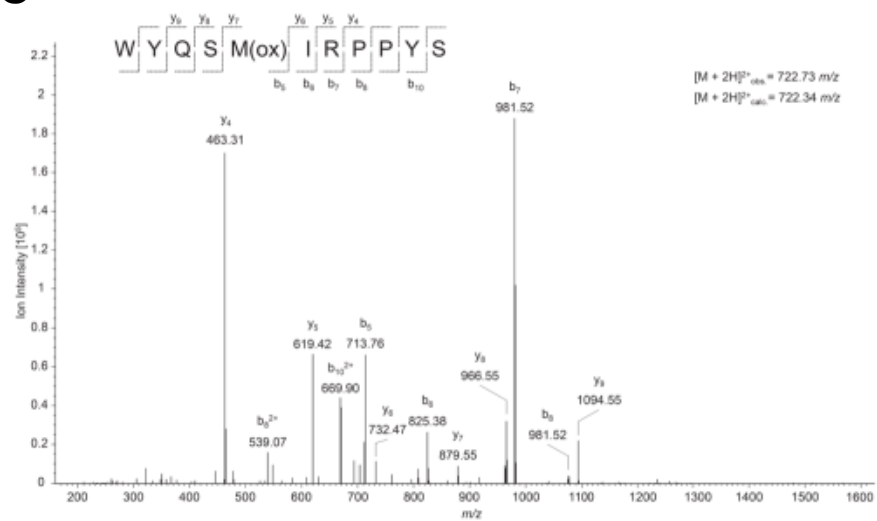
A

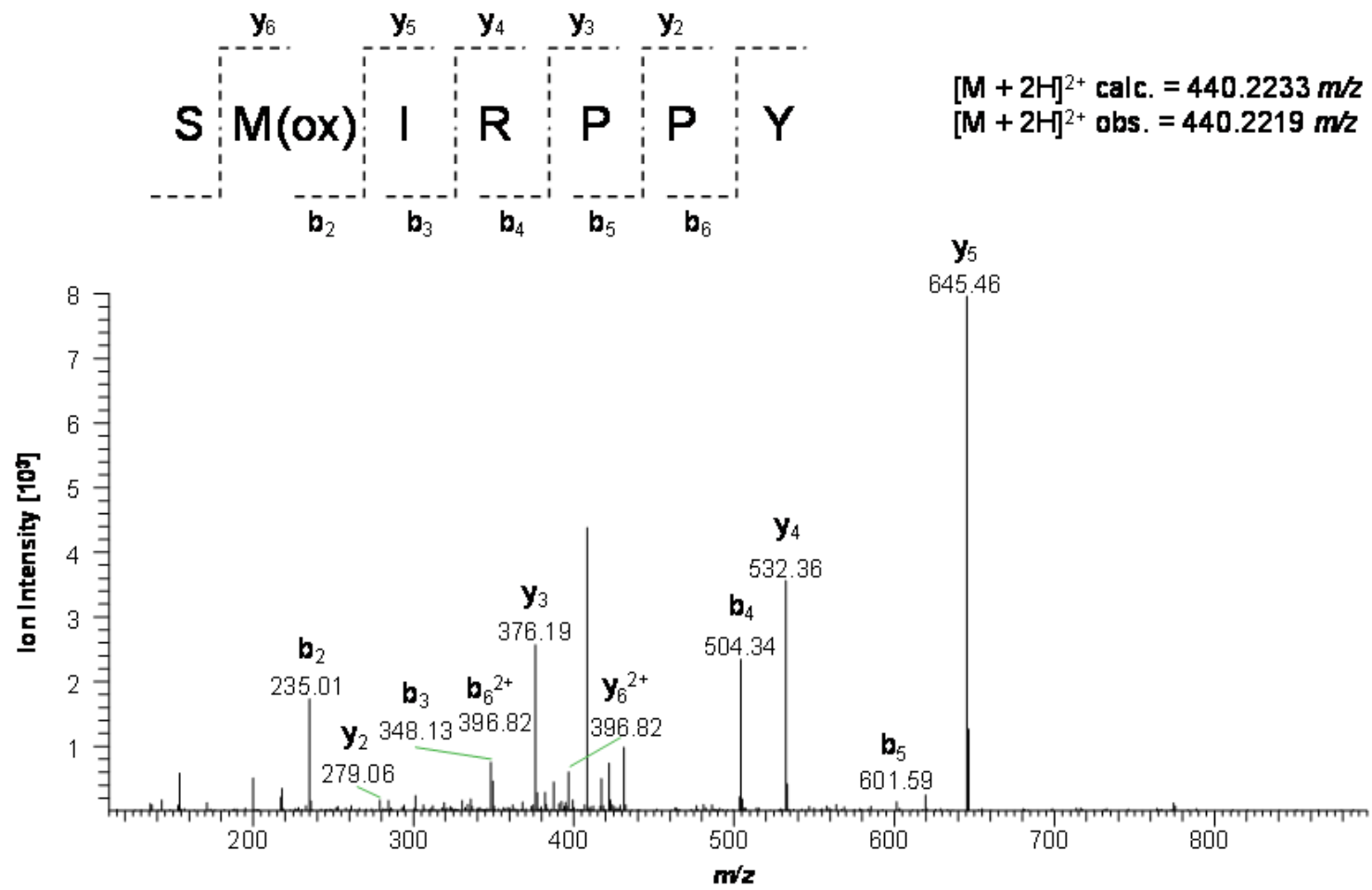


B



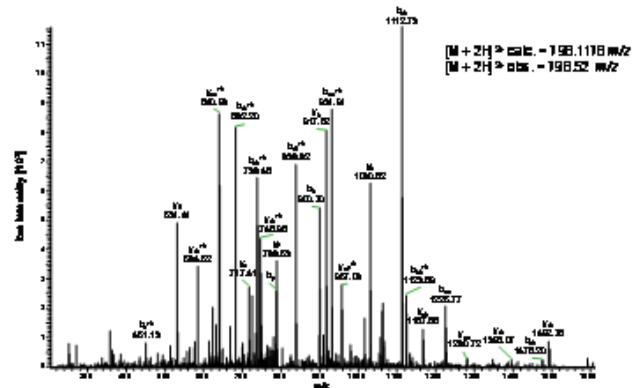
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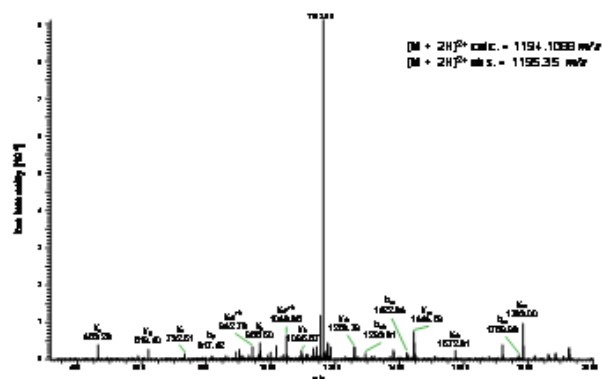
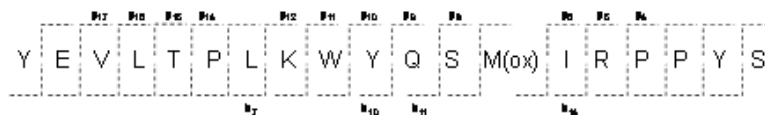


Supplementary Figure 2

A

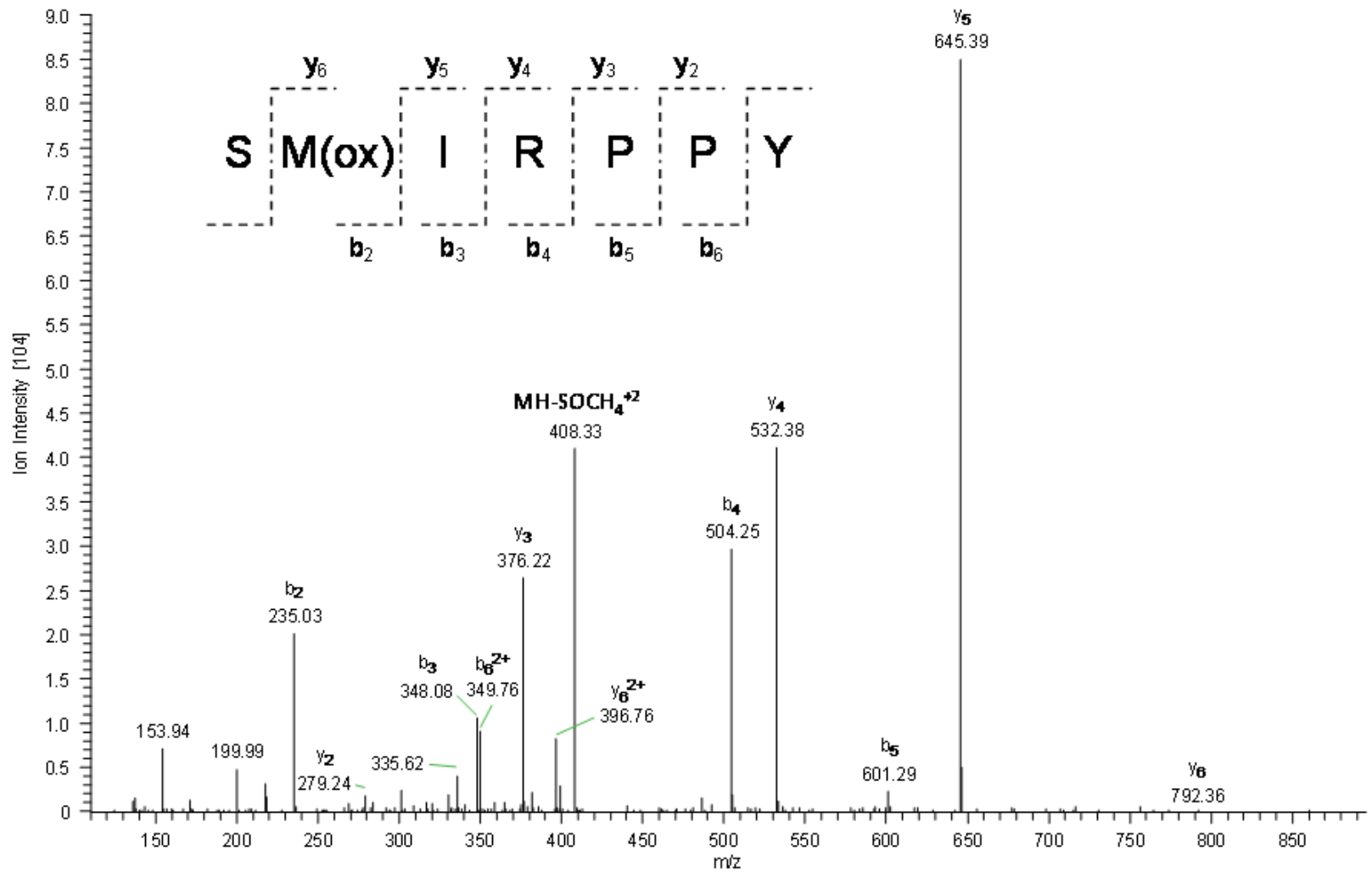


B



Supplementary Figure 3

Gerlach_B8M#4068 RT: 24.24 AV: 1 NL: 9.00E4
T: ITMS + c NSI d w Full ms2 440.22@cid35.00 [110.00-895.00]



Supplementary Figure 4

Supplementary Table 1

Sequence	Count	Amelogenin Isoforms
MPLPPHPG	1	Q99217;Q99217-3;Q99218;Q99218-1
MPLPPHPGHPG	2	
M(ox)PLPPHPGHPG	2	
MPLPPHPGHPGYIN	1	
M(ox)PLPPHPGHPGYINF	4	
LPPHPGHPG	1	
LPPHPGHPGY	2	
LPPHPGHPGYIN	1	
LPPHPGHPGYINF	11	
LPPHPGHPGYINF _S YEVL _T PLK	2	
MPLPPHPGHPGYINF _S YEVL _T PLK	2	
M(ox)PLPPHPGHPGYINF _S YEVL _T PLK	2	
PLPPHPGHPG	1	
PHPGHPGYINF	1	
PHPGHPGYINF	3	
INFSYEVL _T PLK	1	
FSYEVL _T PLK	1	
SYEVL _T PLK	3	
SYEVL _T PLK _W YQ	1	
SYEVL _T PLK _W YQ _S IRPPYP	1	
YEVL _T PLK	2	
YEVL _T PLK _W	1	
YEVL _T PLK _W YQ	1	
YEVL _T PLK _W YQ _S	1	
YEVL _T PLK _W YQ _S IRPPYP	6	
PLK _W YQ _S IRPPYP	2	
LTPLK _W YQ _S IRPPYP	1	
TPLK _W YQ _S IRPPYP	4	
WYQ _S IRPP	1	
WYQ _S IRPPYP	1	
WYQ _S IRPPYPS	3	
WYQ _S IRPPYPSY	2	
WYQ _S IRPPYPSY _G	2	
WYQ _S IRPPYPSY _G YEP	1	
WYQ _S IRPPYPSY _G YEP _M (ox) _G	2	
WYQ _S IRPPYPSY _G YEP _M GGW	2	
WYQ _S IRPPYPSY _G YEP _M (ox) _G GW	3	
SIRPPYPSY	2	
SIRPPYPSY _G YEP _M GGW	3	
SIRPPYPSY _G YEP _M (ox) _G GW	1	
MQPLPPMLPDL _T LEAWP _S TDK	1	
PPM(ox)LPDL _T LEAWP _S TDK	1	
LPDL _T LEAWP _S TDK _T KREEVD	1	
DL _T LEAWP _S TDK	5	
LEAWP _S TDK _T KREEVD	1	
EAWP _S TDK _T KREEVD	1	
P _S TDK _T KREEVD	1	
ILFACLLGAAFAMPVLTPLK _W YQ _S IRPPYP	2	
YEVL _T PLK _W YQ _S _M (ox) _I RPPYS	1	
WYQ _S _M _I RPPY	1	
WYQ _S _M (ox) _I RPPY	1	
WYQ _S _M (ox) _I RPPYS	1	
LRPLPPILPDL _H LEAWP _A TDK	2	

Supplementary Table 2

Sequence	Count	Amelogenin Isoforms
MPLPPHPG	2	Q99217;Q99217-3;Q99218;Q99218-1
M(ox)PLPPHPG	1	
MPLPPHPGH	2	
M(ox)PLPPHPGH	1	
MPLPPHPGHP	3	
MPLPPHPGHPG	5	
M(ox)PLPPHPGHPG	4	
M(ox)PLPPHPGHPGY	1	
MPLPPHPGHPGYIN	4	
M(ox)PLPPHPGHPGYIN	5	
MPLPPHPGHPGYINF	8	
M(ox)PLPPHPGHPGYINF	9	
LPPHPGHP	1	
LPPHPGHPG	6	
LPPHPGHPGY	1	
LPPHPGHPGYIN	3	
LPPHPGHPGYINF	21	
PHPGHPGYINF	7	
PLPPHPGHPG	1	
PPHPGHPGYINF	3	
SYEVLTPK	5	
SYEVLTPKW	2	
YEVLTPK	1	
YEVLTPKW	2	
YEVLTPKWY	2	
YEVLTPKWYQ	4	
SIRPPPSY	5	Q99217;Q99217-2;Q99217-3
SIRPPPSYG	1	
SIRPPPSYGYEP	2	
SIRPPPSYGYEPM	2	
SIRPPPSYGYEPM(ox)	2	
SIRPPPSYGYEPMG	2	
SIRPPPSYGYEPM(ox)G	3	
IRPPPSY	1	
IRPPPSYGYEPMG	1	
AWPSTDKTKREEVD	1	
PSTDKTKREEVD	1	
SM(ox)IRPPY	1	Q99218;Q99218-1

- **Appendices (Supplemental Data)**
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- **Supplementary Table 1.** Peptide sequences identified from amelogenin isoforms from first set of samples (with reductive alkylation and trypsin digest step). Peptides from isoform Y of amelogenin are shown in bold
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- **Supplementary Table 2.** Peptide sequences identified from amelogenin isoforms from second set of samples (simple acid etch). Peptide from isoform Y of amelogenin is shown in bold
-
- **Supplementary Table 3.** Peptide sequences identified from amelogenin isoforms from c. 600 -900 AD samples (simple acid etch)
-
- **Supplementary Figure 1.** Characteristic CID MS/MS spectra of the dimorphic peptides from isoforms X and Y of amelogenin: **A.** WYQSIRPP (from female sample, $[M+2H]^{2+} = 524.15 \text{ m/z}$), **B.** the oxidized form of the WYQSM(ox)IRPPYS (from male sample $[M+2H]^{2+} = 722.73 \text{ m/z}$) and **C.** WYQSMIRPPY peptide (from another male sample, $[M+2H]^{2+} = 671.46 \text{ m/z}$)
-
- **Supplementary Figure 2.** CID MS/MS spectrum of the oxidized peptide from isoform Y of amelogenin: SM(ox)IRPPY (from male sample, $[M+2H]^{2+} = 440.2219 \text{ m/z}$ recorded in orbitrap)
-
- **Supplementary Figure 3.** CID MS/MS spectra of two additional dimorphic peptides from isoform Y of amelogenin identified from male samples: **A.** LRPLPPILPDLHLEAWPATDK and **B.** YEVLTPK WYQSM(ox)IRPPYS
-
- **Supplementary Figure 4.** CID MS/MS spectrum of the oxidized peptide from isoform Y of amelogenin: SM(ox)IRPPY from archaeological male sample, $[M+2H]^{2+} = 440.2230 \text{ m/z}$ recorded in orbitrap)