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Impurity profiling and structure elucidation of phosphoramidite raw materials used for oligonucleotide synthesis

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Solid phase oligonucleotide synthesis

Typical synthesis sequence (followed by purification)





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Phosphoramidite Building Blocks

1. Roy, S.; Caruthers, M. "Synthesis of DNA/RNA and Their Analogs via Phosphoramidite and H-Phosphonate Chemistries" *Molecules*, 2013, 18, 14268-14284. (doi: 10.3390/molecules181114268)

Phosphoramidite impurities can impact the final product





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Incorporated, and may form branched oligomer



2. Kiesman, W.F. et al. "Perspectives on the Designation of Oligonucleotide Starting Materials" Nucleic Acid Therapeutics, 2021, 31, 93-113. (doi: 10.1089/nat.2020.0909)

Case study: 5'-DMT-2'-F-A(bz)-CEP

5'-Dimethoxytrityl-2'-fluoro-benzoyl-adenosine cyanoethyl phosphoramidite



- Obtained material from four different vendors
- Analyze to detect and identify impurities above 0.01% relative abundance using LC/HRAM-MS

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Overview of impurity profiling workflow



LC/MS Setup

Thermo Scientific[™] Vanquish[™] Horizon UHPLC system + Orbitrap Exploris[™] 120 mass spectrometer

- Thermo Scientific[™] Hypersil GOLD[™] C18 column (2.1x100 mm, 1.9 µm)
- Mob. Phases: (A) 10 mM Ammonium acetate, (B) ACN
- **15 min gradient** + 5 min re-equilibration
- UV 200-400 nm
- Full Scan data acquired w/ Polarity Switching
- Resolution Setting 60,000 / 15,000 (MS¹ / MS²)
- *m/z* 200-1200 Da
- Top2 data-dependent MS² with HCD (stepped NCE 10,20,40)





2'-F-A(bz)-CEP samples dissolved in anhydrous ACN at 1 mg/mL to prevent oxidation prior to analysis

Impurity spike-in experiments

Confirming sensitivity of analytical method to detect impurities at 0.1% relative to main component



Impurity spike-in experiments



Analysis of several vendor's materials

5'-DMT-2'-F-A(bz)-CEP – Purity > 99% for all



Analysis of several vendor's materials – detail view



Analysis of several vendor's materials – detail view



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Identity of raw material from MS data

5'-DMT-2'-F-A(bz)-CEP



Structure confirmation of raw material identity by MS²

5'-DMT-2'-F-A(bz)-CEP – automatic fragment structure assignment from *in-silico* prediction



Identification of impurities in Compound Discoverer



1. Compound Selection Thermo Scientific Compound Discoverer software 2-F-A(bz)-CEP (C47 H51 F N7 O7 P) Compounds **Expected + Unexpected Compounds workflows** 2. Dealkylation True Apply Dealkylation Apply Dearylation True Max. # Steps 150 Min. Mass [Da] 3. Transformations **Generate Expected** Dehydration (H2 O ->); Desaturation Phase I Compounds Methylation (H -> C H3) Phase II Chlorination (H -> Cl); Demethylatic Others Max. # Phase II 1 **Create Analog Trace** Detect Compounds 3 Max. # All Steps Find Expected Unbiased Compounds component

Considers dealkylation reactions and other (user-modifiable) transformations



detection

Impurity identification from predicted transformation

Compound **3** - Δ MW = 200 Da (- C₉H₁₇N₂OP)



Only one peak → loss of stereocenter

Confirmation of impurity 3 structure with MS²

$\Delta MW = 200 \text{ Da} (-C_9H_{17}N_2OP) \triangleq CEP$



Compounds 5 + 6 – pair of diastereomers



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Compounds **5** + **6** – pair of diastereomers

Compounds 💎		Compounds per	File Expe	cted Compounds 💡 Expected (Compounds per File		Featu
F	Formula		Calc. MW	Annot. ∆Mass [ppm]	Reference Ion	m/z	RT [min]	MS2
1 🕀	C33 H37	7 F N7 O6 P	677.25252	-0.26	[M+H]+1	678.25981	6.815	

$$\Delta Formula = - C_{14}H_{14}O$$

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Compounds 5 + 6 - pair of diastereomers - Comparison to 5'-DMT-2'-F-A(bz)-CEP



Compounds 🍸		Compounds per	er File Expe		ted Compounds 💡 🛛 Expected (Compounds per File		Featu
F	Formula		Calc.	MW	Annot. ∆Mass [ppm]	Reference Ion	m/z	RT [min]	MS2
1 👳	C33 H37	7 F N7 O6 P	677.2	25252	-0.26	[M+H]+1	678.25981	6.815	



 $-DMT = -C_{21}H_{19}O_2$

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Structure proposal fits observed fragments

MS² spectrum

MS² distinguishes isomeric impurities 21-24

[5'-DMT-2'-F-A(bz)-CEP]-H+CI *m/z* 910.3254



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MS² distinguishes isomeric impurities 21-24



MS² distinguishes isomeric impurities 21-24

5'-DMT-2'-F-A(bz-CI)-CEP and 5'-DMT-CI-2'-F-A(bz)-CEP



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Determination of transformation site allows to trace back to origin of Chlorine substitution

Distinguishing co-eluting impurities with HRAM MS

Peaks 11 and 13 – overlapping MS components



Distinguishing co-eluting impurities with HRAM MS

Peaks 11 and 13 – overlapping MS components – identification based on accurate mass and MS²



Distinguishing co-eluting impurities with HRAM MS

Peaks 11 and 13 - 5'-CEP-2'-F-A(bz)-CEP and 5'-DMT-2'-F-A-CEP





Summary of impurity analysis of 5'-DMT-2'-F-A(bz)-CEP



					-				
Impurity	MS peak (Da)	Formula	Putative	Example impurity	% of total UV peak area by vendor				
impunty			identification	classification	Α	В	С	D	
I	808.2422	$C_{41}H_{38}FN_6O_9P$	-DIPA +O ₂ to P(V) species	Noncritical	0.01%	0.00%	0.00%	0.00%	
Ш	675.2491	$C_{38}H_{34}FN_5O_6$	-CEP	Noncritical	0.00%	0.00%	0.09%	0.00%	
IV	792.2472	C ₄₁ H ₃₈ FN ₆ O ₈ P	-DIPA + OH	Noncritical	0.07%	0.05%	0.07%	0.05%	
V	677.2525	C ₃₃ H ₃₇ FN ₇ O ₆ P	-DMT +bz	Noncritical	0.05%	0.00%	0.00%	0.00%	
VI	891.3518	$\mathrm{C_{47}H_{51}FN_7O_8P}$	Oxidation to P(V) species	Noncritical	0.06%	0.10%	0.13%	0.13%	
VII	822.3306	$C_{44}H_{48}FN_6O_7P$	HP=O phosphite + loss of cyanoethyl group	IP=O phosphite + loss of cyanoethyl group Noncritical 0.01% 0.04%		0.04%	0.04%	0.00%	
VIII	834.2944	C ₄₄ H ₄₄ FN ₆ O ₈ P	Substitution of DIPA with iPrOH	Noncritical	0.06%	0.00%	0.00%	0.00%	
IX	773.3341	$C_{35}H_{50}FN_9O_6P_2$	-DMT +CEP	Critical	0.01%	0.05%	0.07%	0.04%	
Х	771.3307	$C_{40}H_{47}FN_7O_6P$	-bz	Critical				0.04 /0	
XI	779.2753	$C_{45}H_{38}FN_5O_7$	-CEP + bz	Noncritical	0.01%	0.01%	0.01%	0.00%	
XII	861.3414	$C_{46}H_{49}FN_7O_7P$	Demethylation on CEP	Instrument Noncritical 0.00% 0.03%		0.03%	0.00%	0.01%	
XIII	903.3518	$C_{47}H_{55}FN_{3}O_{12}P$	Acetyl-methyl substitution on DMT	Noncritical	0.00%	0.01%	0.00%	0.01%	
XV	992.3551	$C_{50}H_{55}FN_8O_9P_2$	-DIPA +CEP	Critical	0.00%	0.00%	0.03%	0.00%	
XVI	874.3732	$C_{47}H_{52}FN_8O_6P$	-O +NH on bz	on bz Noncritical		0.00%	0.04%	0.00%	
XVII	889.3721	C ₄₈ H ₅₃ FN ₇ O ₇ P	Methylation on DMT	Noncritical	0.09%	0.00%	0.00%	0.00%	
XVIII	909.3177	C47H50CIFN7O7P	Chlorination on bz	Noncritical	0.12%	0.07%	0.05%	0.09%	
XIX	977.3799	$C_{59}H_{52}FN_5O_8$	-CEP +DMT	Noncritical	0.02%	0.01%	0.03%	0.02%	
XX	909.3177	C47H50CIFN7O7P	Chlorination on DMT	Noncritical 0.00%		0.01%	0.01%	0.04%	
XXI	864.3779	$C_{47}H_{54}FN_6O_7P$	-CN +CH ₃ "M-11"	Critical	0.02%	0.00%	0.00%	0.00%	
Р	875.3571	C ₄₇ H ₅₁ FN ₇ O ₇ P	5'-DMT-2'-F- A(bz)CEP	-	99.35%	99.61%	99.44%	99.61%	
				Critical impurity level	0.03%	0.05%	0.10%	0.04%	

For more detail: <u>Application Note AN001949</u>

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Impurity XX

Chemical Formula: C₅₉H₅₂FN₃O₇ Exact Mass: 977.3800 Impurity XIX

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- Phosphoramidites are raw materials for oligonucleotide-derived drug substances, and their impurities need to be controlled in the manufacturing of therapeutic oligonucleotides
- Vanquish UHPLC system coupled with Orbitrap Exploris 120 MS facilitates confident profiling and identification of phosphoramidite impurities
- **Compound Discoverer** software **automates mass spectral annotation** process, allowing users to determine transformation sites
- Fragmentation data allows the distinction of isomeric impurities and to determine the origin of transformations



Thank you

Questions?

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