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## MicroRNA profiling as novel biomarkers for detecting gutter oil using machine learning

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Category	Method	References	Comments
Physical Chemistry	Planar toroidal dipole metamaterial sensor	[15]	These methods typically require specialized detection equipment and are still in developmental stages, resulting in relatively high testing costs. Additionally, conductivity detection is influenced by the refining process of gutter oil, leading to significant fluctuations under deep refining conditions.
	Highly sensitive optical fluid detection with hybrid-waveguide couplers	[16]	
	Raman spectroscopy for identifying polycyclic aromatic hydrocarbons (PAHs)	[17]	
	Conductivity measurement	[56]	5]
Analytical Chemistry	Mass spectrometry and chromatography techniques (e.g., MALDI-MS, UPLC- MS/MS) to detect a variety of substances in gutter oil, including aldehydes, ketones, capsaicin, cholesterol, surfactants, and species- specific fatty acids		<ul> <li>a) The refining process can effectively remove a significant portion of aldehydes and ketones, <u>making direct detection challenging;</u></li> <li>b) Capsaicin levels are highly affected by cooking practices, as certain foods may be cooked without spices, thus making capsaicin <u>detection less reliable;</u></li> <li>c) Cholesterol detection suffers from low sensitivity, and detecting surfactant residues can be influenced by the refining process;</li> <li>d) Mass spectrometry and chromatography require large-scale equipment, leading to high</li> </ul>
Biochemical	Antigen-antibody interactions (detect capsaicin)	[24–26]	Capsaicin detection is still influenced by cooking practices, and antigen-antibody interactions are not reliable enough. Metabolomics is a highly complex and
	Metabolomics	[27,20]	expensive technique.

 Table S1. Reviews of technologies for detecting gutter oil.

miRNA (Accession number)	Label	Sequence
MIR162a (MI0000194)	RT	GTCGTATCCAGTGCAGGGTCCGAGGTATTCGCACTGGATACGACctggat
	F	TCGCTtcgataaacctctgc
MIR168a (MIMAT0001045)	RT	GTCGTATCCAGTGCAGGGTCCGAGGTATTCGCACTGGATACGACgtcccg
	F	TCGCTtcgcttggtgcagat
MIR166 (MIMAT0000635)	RT	GTCGTATCCAGTGCAGGGTCCGAGGTATTCGCACTGGATACGACggggaa
	F	TCGCTtcggaccaggettca
MIR156a (MIMAT0000618)	RT	GTCGTATCCAGTGCAGGGTCCGAGGTATTCGCACTGGATACGACgtgctc
	F	TCGCTtgacagaagagagt
let-7a (MIMAT0000062)	RT	GTCGTATCCAGTGCAGGGTCCGAGGTATTCGCACTGGATACGACaactat
	F	TCGCTtgaggtagtaggttgt
miR-223 (MIMAT0000280)	RT	GTCGTATCCAGTGCAGGGTCCGAGGTATTCGCACTGGATACGACtggggt
	F	TCGCTtgtcagtttgtcaaat
miR-16 (MIMAT0000069)	RT	GTCGTATCCAGTGCAGGGTCCGAGGTATTCGCACTGGATACGACcgccaa
	F	TCGCTtagcagcacgtaaata
Universal Primer	R	GTGCAGGGTCCGAGGT

Table S2. Sequences of primers and probes used for reverse transcription and TaqMan qPCR.

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