1. Genomics Data Management Smart Contract Vulnerabilities

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| Recentrancy in Genomicontraturagement.listRanGenomicHF(uint256) (GenomicsManagement.solH2784-2796):
| State variables written after the call(s):
| RandFTSRandFTSCount| = RandGenomicHF(EasAFTSCount,RandFTSmartContract.t_tokentd,address(mg.sender),SequencingMethodPrice[1],SequencingMethodPrice[3]) (GenomicsManagement.solH2789)
| RandFTSCount+ + (GenomicsManagement.solH2789)
| SecurityOppositValue[RandFTSCount] = sequenceOppositValue[RandFTSCount] = se
```

```
GenomicsDataManagement.refundRawNFTSequencingPayment(uint256, uint256) (GenomicsManagement.sol#2848-2861) uses timestamp for comparisons

__require(bool_string)(block.timestamp > SequencingRequestStartTIme[Raw.RawNFTId][_requestnumber] + SequencingPeriod, The time window for sequencing is still openned) (GenomicsManagement.sol#2853)

Bangerous comparisons:

__require(bool_string)(block.timestamp > SequencingRequestStartTIme[Raw.RawNFTId][_requestnumber] + SequencingPeriod, The time window for sequencing is still openned) (GenomicsManagement.sol#2864-2874) uses timestamp for comparisons

Dangerous comparisons:

__require(bool_string)(block.timestamp > SequencingRequestStartTime[Raw.RawNFTId][_requestnumber] + SequencingPeriod + InitialFullAccessGrantingPeriod, The time window for initial full acces is still openned) (GenomicsManagement.sol#2864)

GenomicsDataManagement.ConfirmMawNFTSequencing(uint256, uint256) (GenomicsManagement.sol#2894-2999) uses timestamp for comparisons

__nequire(bool_string)(block.timestamp <= SequencingRequestStartTime[Raw.RawNFTId][_requestnumber] + SequencingPeriod, The sequencing time window is already closed) (GenomicsManagement.sol#2913-2929) uses timestamp for comparisons

__nequire(bool_string)(block.timestamp <= SequencingRequestStartTime[Raw.RawNFTId][_requestnumber] + SequencingPeriod + InitialFullAccessGrantingPeriod, The Initial Full Access Granting Period time window is already closed) (GenomicsManagement.sol#2913)

GenomicsDataManagement.Jatamagement.Sol#29130)

GenomicsDataManagement.Jatamagement.Sol#29130 (GenomicsManagement.sol#293-3033) uses timestamp for comparisons

__require(bool_string)(block.timestamp <= SequencingRequestStartTingTime[Sequenced.SeqNFTId][] [_requestnumber] + SequencingPeriod + InitialFullAccessGrantingPeriod, The Initial Full Access Granting Period + InitialFullAccessGrantingPeriod, The Initial Full Access Granting Period + InitialFullAccessGrantingPeriod, The Initial Full Access Granting Period + InitialFullAccessGrantingPeriod, The InitialFullAcc
```

```
Renetromy in Genomicalizationagement. Confirm@wifeTSequencing(intr256_uint256) (Genomicalizagement, sol#2004):

terenal calls:

- address(mag. sender). transfer (SequencingfethodPire[RequestedSequencingfethod[Raw,RawNTId][_requestrumber]]) (Genomicalizagement, sol#2003)

State variables sertice after the call(s):

- Sequencing(sequencing). (Faut.RawFITd] - rel (Genomicalizagement, sol#2005)

rever senticed greater the call(s):

- Sequencing(sequencing). (Faut.RawFITd] = rel (Genomicalizagement, sol#2005)

Renetranny. In Genomicalizationagement. InitialFull_(requestrumber] - true (Genomicalizationagement, sol#2005)

Renetranny. In Genomicalizationagement. InitialFull_(Access/teromicalizagement). (Genomicalizagement, sol#2004)

Renetranny. In Genomicalizationagement. InitialFull_(Access/teromicalizagement). (Genomicalizagement). (Genomicalizagement
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2. Raw Genomic Data NFT Smart Contract

3. Sequenced Genomic Data NFT Smart Contract