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| **Addiitonal file 3: Table S3a** **Correlation of PODXL score between whole tissue sections and corresponding TMA cores from resected primary tumors** |
|  | **PODXL score in TMA cores** |
| **PODXL score in whole tissue sections** | 0 | 1 | 2 | 3 | 4 | Correlation (*τ*b)p |
| 0 | 936.0% | 00.0% | 00.0% | 00.0% | 00.0% | 0.944**<0.001** |
| 1 | 14.0% | 416.0% | 00.0% | 00.0% | 00.0% |
| 2 | 00.0% | 14.0% | 416.0% | 00.0% | 00.0% |
| 3 | 00.0% | 00.0% | 00.0% | 416.0% | 14.0% |
| 4 | 00.0% | 00.0% | 00.0% | 00.0% | 14.0% |

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| **Addiitonal file 3: Table S3b** **Correlation of PODXL expression (trichotomized/dichotomized) between whole tissue sections and corresponding TMA cores from resected primary tumors** |
|  | **PODXL (trichotomized) in TMA cores** |  | **PODXL (dichotomized) in TMA cores** |
| **PODXL (trichotomized) in whole tissue sections** | Negative | Low | High | Correlation (*τ*b)p | **PODXL (dichotomized) in whole tissue sections** | Negative | Positive | Correlation (*τ*b)p |
| Negative | 936.0% | 00.0% | 00.0% | 0.956**<0.001** | Negative | 936.0% | 00.0% | 0.919**<0.001** |
| Low | 14.0% | 936.0% | 00.0% |
| Positive | 14.0% | 1560.0% |
| High | 00.0% | 00.0% | 624.0% |