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2	Protocol
3	Main inclusion criteria:
4	Age≥18, Pathologically identified non-small cell lung cancer
5	Adenocarcinoma, No-drive gene mutations (with EGFR, ALK, ROS1 included detected by next
6	generation sequencing, NGS)
7	Squamous cell carcinoma, there is no need to perform gene detection but will be retrospectively
8	detected by NGS
9	Progression disease after second line treatment
10	Patients have not been administrated with Vinorelbine or apatinib
11	PS score ≤ 2
12	Estimated survival time more than 3 months
13	
14	Main exclusion criteria:
15	Patients received treatment of apatinib or Vinorelbine before
16	EGFR, ALK or ROS1 mutation positive for Lung Adenocarcinoma patients
17	EGFR, ALK or ROS1 mutation positive for Squamous cell carcinoma if the patients have
18	performed gene detection by NGS
19	Patients with contraindication of chemotherapy
20	Pregnant or breast feeding women
21	
22	Outcome Measures:
23	Primary Outcome: Objective Response Rate (ORR)
24	ORR means the best response status during the whole treatment line.
25	Secondary Outcome Measures: Progression free survival (PFS), OS (Overall survival) and safety
26	
27	Sample size calculation:
28	We have chosen a fixed sample size of 30
29	
30	Statistical analysis:
31	SPSS (version 22.0) was used for statistical analyses. Proportion of responders were calculated

with the 95% confidence intervals (CI) using the Clopper-Pearson method. Kaplan-Meier method

was used to estimate the median duration of response and PFS with corresponding 95% CIs.

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